

RESEARCH ARTICLE

Plastic Degrading Enzymes and Microbes

Dipsikha Bordoloi¹, Nishita Acharjya¹ and Biswa Prasun Chatterji^{2*}

¹B.Sc Final Year Student, Department of Biotechnology, Assam down town University, Assam, India. ²Professor, Department of Biotechnology, Assam down town University, Assam, India.

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*Address for Correspondence Biswa Prasun Chatterji

Professor, Department of Biotechnology, Assam down town University, Assam, India. Email: biswaprasun@gmail.com



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ABSTRACT

Plastic pollution poses a significant environment challenge, and the research for sustainable solutions has led to the discovery of plastic -degrading enzymes. This article provides a comprehensive overview of the recent advances in the field of plastic degrading enzymes, offering insights into the mechanisms, substrates, and potential applications of these remarkable biocatalysts. We examine the diversity of enzymes capable of degrading various types of plastics, including polyethylene, and PET and delve into the genetic and biochemical basis for their efficacy. Furthermore, we discuss the environmental impact of these enzymes, their industrial applications and the challenges and the opportunities that lie ahead in harnessing them for large -scale plastic waste remediation.

Keywords: enzymes, microbes, biodegradation, plastic

INTRODUCTION

Plastic is one of the biggest concern that humanity is facing today. Plastic pollution has become one of the major environmental problem .The widespread use of plastic materials has led to an accumulation of plastic waste in our oceans, landfills, and ecosystems. The world plastic production has reached 350 million tons till 2017(, Demand Plastics-the Facts 2018 An Analysis of European Plastics Production and Waste Data, n.d.) Plastic is a synthetic material made from polymers, which are long chains of molecules. Polymers are created by linking together smaller molecules called monomers through a process known as polymerization. The specific composition of plastics can vary widely, as there are many different types of plastics with different and unique properties. Polyethylene(PE), Polyethylene terephthalate(PU), Polystyrene (PS), polypropylene(PP), and polyvinyl chloride (PVC) are the main types of plastic which accumulation in the environment is a major concern for environment and human health.(Hassan et al., 2022)These synthetic plastics have been separated into two groups based on the degradation





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pathways, plastics with a carbon-carbon backbone and plastics with heteroatoms in the main chain .(Mohanan et al., 2020). The sole component of the backbone of PE,PS,PP, and PVC polymers is carbon atoms. Microplastics (MPs) are plastic particles smaller than 5 mm that are produced during plastic breakdown and may have ecotoxicological effects. (Zhang et al., 2017) Inhaled fibrous particulates have the potential to remain in the lungs and, when combined with other contaminants like plasticizers and dyes, can have harmful consequences on health, such as mutagenicity and cancer. It is acknowledged that burning plastic garbage can permanently remove it. Nonetheless, unburned debris is still present in the bottom ash, which is a solid residue left behind from incinerators that can burn up to 360–102,000 microplastic particles per metric ton. One possible source of MPs released into the environment is this bottom ash. According to reports, plastic pieces known as nanoplastics that are smaller than 100 nm. In general, commonly used plastics donot break down significantly in the environment when released. This should come as no surprise given that many polymers remarkably high stability and durability are among the main factors contributing to their widespread application and acceptance. Plastic breaks down in the environment through four different mechanisms. Photodegradation, thermooxidative degradation, hydrolytic degradation and microorganism – mediated biodegradation. (Webb et al., 2013)

POLYMER DEGRADATION

Polymer degradation can happen through two separate mechanisms: bulk erosion and surface erosion. The process of surface erosion is restricted to the device's surface. Overtime, the polymeric device will shrink in size without losing its bulk integrity. This kind of erosion affects poly(ortho-esters)and polyanhydrides. On the hand, bulk erosion occurs when water enters the device more quickly than the polymer is changed into components that are soluble in water, causing erosion all the way through the device. There are two steps to this process, the first step – water permeates the majority of the apparatus, primarily cleavage the chemical bonds in the amorphous phase that are hydrolytically unstable and breaking lengthy polymer chains into shorter segments that are eventually soluble in water. Enzymatic assault of the fragments takes place in the second phase. PGA and I-PLA are examples of semicrystalline polymers that exhibit this kind of erosion. The chemical stability of polymer backbone, the hydrophilic/hydrophobic balance of the repeating units, the polymer's shape (semicrystalline/ amorphous), and its molecular weight and molecular weight distribution are the primary determinants of erosion rate. Since ester bonds serve as links between the repeating units of PGA, PLA, and copolymers PLG, they should all erode at the same rate if the chemical stability of the polymer backbone is the only factor influencing the rate of erosion. However, the ability of the water molecules is influenced by the hydrophilic versus hydrophobic nature of the repeating units. As a result, PGA erodes far more quickly than PLA, which is more hydrophobic. Furthermore, the polymer's shape is quite significant. Water is unable to enter the crystalline domains in the crystalline form due to the well- organized and tightly packed polymer chains. As a result, backbone hydrolysis typically occurs on the surface of crystalline domains and in a amorphous regions. The rate of erosion of I-PLA and d,I-PLA serves as an illustration of this process. Despite having the same degree of hydrophobic and backbone links as I-PLA, d,I-PLA breaks down significantly more quickly than I-PLA because I-PLA is an amorphous polymer whereas, the stereo regular polymer

TYPES OF DEGRADATION

Phtooxidative degradation

The main cause of damage, to polymer is light. This process begins when light is absorbed leading to degradation through photodegradation and photooxidation. Synthetic polymers are particularly susceptible, to degradation caused by ultraviolet (UV)radiation. The lifespan of materials used in application is influenced by this process (Riinby,1989) During photo irradiation the soft segments of polymers where degradation occurs can produce ester, aldehyde, propyl and formate groups. UV radiation easily breaks C-C bonds in the polymers.

Ozone degradation

is semicrystalline.

Polymeric degradation is caused by the presence of ozone in the environment. Polymer exhibits a longer half -life in the absence of oxidative reactions. Despite being extremely rare in the atmosphere, ozone has a significant impact on





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polymers. Polymeric materials undergo degradation due to the generation of reactive oxygen species. These ROS are created when the molecular weight of polymers decreases and their mechanical and electrical properties alter. Different kinds of carbonyl and unsaturated carbonyl compounds are generated when polymers are subjected to ozone. These goods are made from aromatic carbonyl, lactones, ketones and esters.

Hydrolytic degradation

The process by which plastic materials disintegrate into smaller components when water is present is known as hydrolytic degradation of plastic. Water molecules usually cause the breaking of chemical bondswithin the polymer structure during this breakdown process. Numerous plastic types are prone to hydrolytic degradation, with the degree of deterioration, with the degree of deterioration contingent upon variable likes temperature, catalyst presence, and polymer makeup.

Biodegradation

Biodegradation is the term used to describe any physical or chemical change in a material brought about by microorganisms. Bacteria, actinomycetes, and fungi are among the microorganisms that break down both natural and manmade plastics.(Cai et al., 2023) There are mainly two types of biodegradations aerobic and anaerobic biodegradation. Using oxygen as an electron acceptor bacteria break down large organic compounds into smaller ones in aerobic type of biodegradation. In this form of degradation this method yields water and carbondioxide as biproducts.

Carbon plastic +O₂ CO₂+H₂O+carbon residual

The breakdown of organic pollutants by microorganisms in the absence of oxygen is known as anaerobic biodegradation. At hazardous waste sites, it plays a significant role in the natural attenuation of Pollutant. Certain anaerobic bacteria break down organic compounds into smaller ones by using carbon dioxide, nitrate, sulphate, iron, manganese, and other elements as their electron acceptors. Since , polymer molecules are lengthy and not soluble in water, microorganisms cannot carry the polymers directly through their outer cell membranes and into the cell where the majority of metabolic reaction occur. Microbes evolve a technique through which they secrete extra cellular enzymes that depolymerize the polymer outside the cells in order to use such material as a carbon and energy source.

Mechanism of biodegradation

The specific mechanism of biodegradation can vary depending on the type of material being degraded and the microorganism involved.(Saritha et al., 2021)

- 1. Recognition and uptake-Microorganisms recognize and ingest the organic material to be degraded. This is often facilitated by specialized enzymes or receptors on the cell surface.
- Extracellular enzymatic action-some microorganisms release enzymes into their surrounding environment.
 These enzymes break down complex organic molecules, like carbohydrates, proteins, and lipids, into smaller, more manageable compounds. For example, amylase, breaks down starch and protease breakdown proteins.
- 3. Transport of breakdown products-The smaller compounds produced by enzymatic action are transported into the microbial cell. this is essential step as it allows the microorganism to utilize these compounds as a source of energy and nutrients.
- 4. Intracellular metabolism-Inside the microbial cell, these compounds undergo further degradation through various metabolic pathways. microorganisms extract energy and building blocks for their own growth and reproduction.
- 5. Final metabolic products-As the compounds continue to break down, they may be converted into CO2and H2O, and other simpler molecules. These end products are usually non-toxic and can be released into the environment without harm.

Bacteria- Bacteria play a significant role in the biodegradation of plastics, particularly in the degradation of PE and PP. Bacteria first need to colonize the plastic surface. They can attach themselves to the plastic through the secretion of extracellular polymeric substances, which creates a biofilm. This biofilm provides a suitable environment for





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bacterial growth and enzyme production. Once attached, bacteria is secrete enzymes such as lipases, esterase, and proteases. These enzymes hydrolyze the polymer chains in plastics into smaller, more easily digestible fragments. Bacteria then metabolize the smaller plastic fragments, using them as a source of carbon and energy. (Atanasova et al., 2021)As the polymer chains are broken down further, the plastic becomes more susceptible to biodegradation. The bacteria can continue to feed on the plastic until it is completely broken down into toxic byproducts. The fungisome species of fungi like aspergillus and penicillium, have been found to have the ability to degrade low-density polyethylene. (Srikanth et al., 2022)Fungi use enzymatic processes to degrade plastic. Enzymes like esterases, lipases, and peroxidase can cleave the chemical bonds in plastic polymers. Microbial polymers degradation includes biofragmentation, biodeterioration, mineralization, and assimilation. Biodeterioration is the procedure that modifies the chemical and physical properties of plastics as well as their surface. (Schmidt, 2016) The mix and structure of polymers determine all chemical and structural alterations. Following biodeterioration plastic polymers are subjected to an enzymatic process known as bio-fragmentation. Oxygenases, which are primarily found in bacteria, are enzymes that can break down oxygen molecules when they are introduced to carbon chains, this process produces less hazardous compounds like alcohol and peroxyl. Moreover, lipases, esterase's, and endopeptidases, for amide groups, catalyze the transformation process of carboxylic groups. Mineral Deposition Via cell membranes, plastic polymers generated during the bio-fragmentation process enter microbial cells. Large monomers are unable to get through the cell membrane and remain outside. The energy is produced by oxidizing the tiny monomers that migrated within the cells. Eventually, this energy is used to produce biomass. In the assimilation process, atoms are integrated in the microbial cells for complete degradation.

Enzymatic degradation of plastic

Degrading plastic with microbial enzyme is a particularly challenging process since the carbon-carbon backbone of plastic lacks hydrolysable groups. The fast stage that is accomplished by the interaction of biotic and abiotic variables is the lowering of molecular weight. Microbial enzymes can readily attack the carbonyl group of a polymer when exposed to UV radiation Several enzymes are employed to break down polymers, including laccase, urease, lipase, protease, and manganese-dependent enzymes (which break down lignin).(Kaushal et al., 2021)Polyethylene (PE) can be broken down by thermostated laccase after 48 hours of incubation at 37°c *Ideonella sakaiensis*, a bacteria that can degrade polyethylene terephthalate, a popular plastic used in bottles and containers, was found by researchers in 2016. PETase and MHETase are the two enzymes that this bacterium produces. PET is broken down into smaller monomers by PETase, and the monomers are further broken down by METase. These enzyme support PET plastic recycling. Scientist have been attempting to modify these enzymes in order to increase their capacity to break down plastic. A biodegradable plastic made from renewable resources like cornstarch is called polylactic acid. however, PLA waste might not break down quickly if it finds itself in unsuitable surrounding, which will facilitate PLA breakdown in composting sites and natural settings.

CONCLUSION

This timeline marks the age of inventive and advanced practices where the world of conservation and engineering go hand in hand. The study of plastic degrading enzymes and their interaction with the microbes has opened up exciting possibilities for addressing the global plastic pollution crisis. It is very important to highlight the remarkable progress in our understanding of these enzymatic and microbial processes , shedding light on their potential for sustainable plastic degradation. As we move forward , it is evident that further research and collaboration among scientists, engineers, and environmentalists will be essential to harness this knowledge effectively. The development of innovative biotechnological solutions and the implementation of eco friendly practices hold the promise of mitigating the detrimental impacts of plastic waste on our planet. The future of plastic degradation appears brighter , thanks to the synergy between this remarkable enzymes and natures microorganisms, offering hope for a cleaner , more sustainable future.





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Table1-classification of polymers with its lifespan and density(Ru et al., 2020)

POLYMERS	DENSITY	LIFE SPAN
PET	1.35	450
HDPE	0.94-0.97	>600
LDPE	0.91-0.93	10-600
PS	1.03-1.09	50-80
PP	0.90-0.91	10-600
PVC	1.35-1.45	50-150

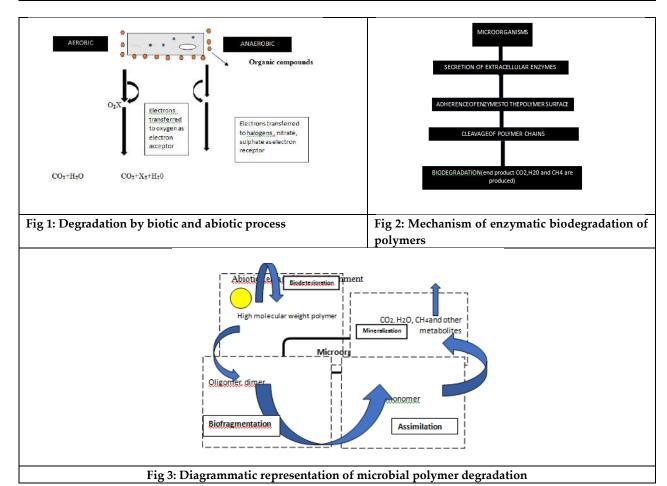




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Table2- list of microbes and its enzyme involved in plastic degradation

MICROBES	ENZYMES	PLASTIC TYPES	MICROBIAL SOURCE	EFFECIENCY OF DEGRADATION
Bacillus cereus	phospholipases	polyethylene	Dumping soil	2.40-7.20
Pseudomonas putida	lipase	Milk cover	Garden soil	75.30
Streptomyces sp	proteases	LDPE	Garbage soil	46.70
Pseudomonas sp	lipase	Polyethylene	Textile effluent	39.70-19.60
Aspergillus glaucus	Pectinase	Plastic and polythene	Mangrove soils	20.80-7.26
Micrococcus luteus	glycosidase	Plastic cups	Forest soils	38.00







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RESEARCH ARTICLE

A Comprehensive Survey of Image Encryption Techniques Enhanced By **Digital Signatures**

C.Sasthi Kumar* and N.Kanagaraj

Assistant Professor, Department of Computer Science, Karpagam Academy of Higher Education, Coimbatore, Tamil Nadu, india.

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*Address for Correspondence

C Sasthi Kumar

Assistant Professor,

Department of Computer Science,

Karpagam Academy of Higher Education,

Coimbatore, Tamil Nadu, india.

Email: sasthikumar.chandrasekar@kahedu.edu.in



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ABSTRACT

As the digital era progresses, the safe transmission and storage of visual data, especially photographs, has become critical. To protect the security and integrity of these visual assets, image encryption algorithms have been created. This survey study investigates the new convergence of picture encryption and digital signature technologies, presenting a novel solution that provides both security and visual meaningfulness. Traditional picture encryption approaches generally rely on data obfuscation through mathematical transformations, which often renders decrypted images aesthetically unintelligible. In contrast, our suggested technique uses digital signatures to improve security while keeping the significance of the visual information. This survey study analyses and analyzes the essential components of this unique algorithm, emphasizing its merits and applications. The algorithm's fundamental components include a hybrid encryption approach that combines symmetric and asymmetric cryptographic algorithms. This article reviews 26 research papers for image encryption techniques and explores the potential of computer-assisted methods for image encryption techniques and staging. Digital signatures are used to authenticate the integrity and validity of the visual data while also guaranteeing that the decrypted pictures are visually understandable. The survey article examines the algorithm's mathematical underpinnings, encryption and decryption operations, and the incorporation of digital signatures into the picture encryption workflow.

Keywords: Digital Signatures, Data Integrity, Image Encryption, Visual Data Security





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INTRODUCTION

In an era marked by extensive usage of digital photography and broad sharing of visual material, preserving the privacy and security of these photographs has become more important [1-3]. The need for strong picture encryption solutions has never been more critical, given the increased danger of illegal access, interception, and data breaches. Visually meaningful picture encryption is a cutting-edge way of protecting sensitive visual information while keeping the image's core structure and visual coherence [4-8]. Traditional picture encryption algorithms often rely only on scrambling the pixel values, leaving the encrypted image unreadable to the human eye [9]. While efficient for data security, these approaches often produce aesthetically disordered and unintelligible pictures, making them unsuitable for applications where keeping the visual sense of the material is critical [10]. This constraint has motivated the development of visually meaningful picture encryption as a solution that achieves a careful balance between security and visual comprehension [11]. Visually meaningful picture encryption approaches strive to encrypt visual material in such a manner that authorized users may perceive and interpret it while making it unintelligible to unauthorized persons or hostile actors [12]. To accomplish both security and meaningful visual representation, this unique strategy integrates powerful cryptography algorithms with image processing techniques [13-16]. The significance of visually meaningful picture encryption extends across several disciplines. Medical photographs including sensitive patient information, for example, must be securely transferred and kept without affecting the diagnostic usefulness of the images [17-19]. Protecting intellectual data and personal images against illegal access is critical in the era of multimedia communication and content sharing [20-23]. We will explore into the fundamental ideas, methodologies, and applications of visually meaningful picture encryption in this research. We'll look at how cryptographic approaches are used with image processing to create a careful balance between picture security and visual information preservation. Furthermore, we will investigate real-world use cases, problems, and future possibilities of visually meaningful picture encryption, shining light on its importance in our increasingly digitized and linked world [24-26].

Background Study

Survey on Image Encryption Techniques

Abbasi, S. F. et al. (2018) the author offers an innovative and effective method of image encryption. First, a change was performed by randomly rearranging the image's pixels using a tangled web of logistic chaotic maps. XORing the random map with the jumbled image improves the security of the system. Gray S-Box was then used to do a substitution on the resulting perverted image. Finally, the encrypted image was transformed into something comprehensible using lifting wavelet transform. The number of possible assaults on the cipher-text image that was produced was lower since it was not a random or noise-like image. Armijo-Correa, J. et al. (2020) the author offer a new, more visually intuitive method of encrypting images. In specifically, the author have enhanced some steps of a visually meaningful encrypted image (VMEI) system, including: To prevent the resulting VMEI image's pixel values from going crazy, a threshold operation was first investigated and tested to identify a suitable dynamic range for the source images. This method also aids in preserving a high percentage of original pixels, which was essential for preserving the high quality of the final VMEI image. The author used a well-known S-box based on a cellular automaton approach to randomly reorganize the data in the new image to get diffusion characteristics throughout the embedded information.

Bai, S. et al. (2020) Current image encryption techniques convert encrypted images to meaningless jumbles of random data, making them vulnerable to attack. The data amount may have grown substantially, or the calculation may have become quite complicated. To address these limitations, these authors research proposes a new kind of aesthetically meaningful image encryption technique that uses fractal graph formation. The suggested method transforms basic images into substantial fractal sceneries or fractal plants with a low time complexity, specific anti-compression performance, and high security performance. Chai, X.et al. (2017) these authors' research presents a compressive sensing-based visual security system for encrypting images. The original image was encrypted using compressive sensing after being transformed into a coefficient matrix through discrete wavelet transform. Then, a zigzag path





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related to the original plain image was used to jumble it. Second, the author produces a cipher image that was both cryptographically and aesthetically secure by embedding the cipher image within a carrier image. The proposed encryption method has three advantages over the state-of-the-art encryption techniques. It has a high security level, can achieve image data security and appearance security simultaneously, and the final cipher image has the same size as the plain image, so it transmits and stores quickly over the Internet and needs no extra transmission bandwidth or memory, but it was very sensitive to the plain image, the parameters used in the zigzag confusion process, and the generation of the chaos-based measurement.

Chai, X. et al. (2020) the author offer a new, aesthetically meaningful, and efficient double-color picture encryption scheme based on 2D compressive sensing (CS) and an embedding approach. The author uses the 2D CS theory to shrink the regular pictures and get two-dimensional measurements of the originals. The author start with raw measurement value matrices, scramble them using index sequences generated by a 6D hyper chaotic system, and then integrate the resulting cipher image into a color carrier image. In 2D CS, the Kronecker product (KP) and singular value decomposition (SVD) were combined to create the measurement matrices, and the feature parameters of the carrier image were employed to govern the embedding process of the compressed cipher pictures. The LSS and 6D hyper chaotic system's starting settings were derived from the plain pictures' SHA 256 and feature parameters. The plain image parameters were included into the carrier image, and the carrier image parameters may be obtained from the corresponding cipher picture with little additional transmission and storage. The suggested encryption method has been shown to be safe against noise assaults, occlusion attacks, known-plaintext attacks, and chosen-plaintext attacks via simulations and performance assessments. It also has a wide key space and high key sensitivity during decryption. Dolendro Singh, L., & Manglem Singh, K. (2018) these authors research article proposes a method for creating multigame encryption schemes that have some kind of aesthetic value. A multiple encrypted picture was created by embedding data from numerous cipher images into the relatively unimportant actual data of a scrambled host image and then unscrambling the image. The suggested approach offers a greater potential to incorporate more data with nearly no damage of the host picture, as shown by comparison with current visually meaningful encryption systems. Different conventional assaults, such as the salt and pepper noise attack and the occlusion attack, can't break the suggested system.

Survey on Digital Signatures

F. Jie *et al.* (2019) the author offer a compressive sensing and Integer wavelet transform (IWT)-based visual security picture encryption system. After the image has been encrypted using compressive sensing, it was further protected using zigzag confusion and a chaotic map. In addition, it was possible to minimize the quantity of data used in the embedding procedure. Second, the carrier image's three color channels were subjected to IWT, and the resulting coefficient matrices were used to simultaneously embed the picture into the matrices. Inverse IWT may be used to produce the safe-looking picture. The following benefits of the encryption system exist as compared to the current encryption method. First, it has excellent camouflage and was not easily noticeable, allowing you to keep your image's look secure. Second, the encryption method does not rely on the carrier image, and the quality of the reconstructed picture was consistent with that of the original photos. The system employs binary integers in the embedding process, thus even though an image was embedded in the paper; it may encrypt any information that can be converted to binary.

H. R. Vanamala and D. Nandur (2019) Here, an evolutionary algorithm was employed to encrypt data in a way that makes sense visually during the pre-encryption phase. Two chaotic maps, a genetic algorithm, and some visual manipulations were used in the essay to cover up the sensitive image. It was possible to use a genetic algorithm to find the optimal encrypted image. Lowest correlation coefficient indicates optimal encrypted image. The algorithm's vulnerability to statistical assaults, differential attacks, and Bruce force has been shown. Huo, D. *et al.* (2021) Two distinct phases, pre-encryption using dimensional compressive sensing (2DCS) and embedding with IWT, make up the visually safe picture encryption system proposed in this research. Pre-encryption protects the original image's data by replacing it with a hidden image produced by a combination of random grayscale alteration, random scrambling, and 2DCS sampling. First, the carrier image was decomposed into a coefficient matrix via IWT, then the





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secret image was embedded into the lowest two bits of the coefficient matrix, and finally the matrix was transformed into the space domain via inverse IWT to obtain an image that looks like the carrier image while still protecting the plain image. Two-dimensional concatenated substitution (2DCS) allows for faster encryption and decryption without requiring more transmission bandwidth or storage space for the plain picture.

M. Liu et al. (2021) the author offer a method for encrypting color images that has both visual meaning and high performance in areas like embedding and reconstruction. To begin, the picture was compressed using compressive sensing technique, which drastically lowers the overall processing cost. Second, the secret picture's security was bolstered by the employment of chaotic sequences to further confound the compressed image, turning it into an indecipherable blur of noise. The encrypted picture was placed inside a visually relevant carrier image to better secure the security of image information. In order to make it more difficult for an attacker to read the plain image or the noise-like cipher image, the former was masked within the latter, creating a visually meaningful cipher picture. The peak signal noise ratio (PSNR) tests further show that the proposed strategy results in a more satisfying visual experience. P. Ping et al. (2019) Compressive sensing and reversible color alteration were proposed as novel methods for image encryption in this study. During the CS stage, it was possible to use completely arbitrary measurement matrices to evaluate each column of the image signal separately. Multiple hidden versions of the same picture may be created using the same secret key if the random numbers were varied. Thanks to CS at the embedding step, the size of the carrier image was decreased in comparison to the size of the plain picture when the block pairing and replacement approach was utilized, meaning that no more bandwidth was needed for transmission. The security analysis and simulation findings show that the scheme was robust against common attacks, has a large key space, and has high key sensitivity.

Survey on Digital Signatures in Visual Data Security

Ponuma, R. et al. (2019) the author present a multiple image compression-encryption technique that utilizes chaotic measurement matrices and compressive sensing. The experimental findings showed that the system outperformed competing approaches when it came to picture reconstruction. An image-dependent key was used to increase protection against a predetermined plain text attack. The technology was also resistant to outside influences like noise and occlusion. The correlation study demonstrates that the suggested strategy was resistant to statistical assaults. The effectiveness and safety of the proposed approach have been shown via simulation. Priya, S., & Santhi, B. (2019) the medical image and the associated patient data were both protected via watermarking. Normal medical image encryption yields an encrypted image with noise or texture that was unsuitable for transmitting the watermarked medical image. The author suggested using an encrypted picture with symbolic value. To create a watermarked medical picture, Electronic patient record (EPR) data was incorporated inside the image utilizing medical image watermarking. The fingerprint was then used to visually encrypt this watermarked medical picture using IWT. The fingerprint image was validated and decoded from the wavelet coefficients on the extraction side. If the fingerprint was a good match, the medical image may be rebuilt and the Extended Producer's Responsibility (EPR) extracted for use in diagnosis. According to the findings of the experiments, the suggested approach was successful in creating an encrypted picture with visual significance.

V. Himthani *et al.* (2022) multiple types of Meaningful Encrypted Image (MEI) techniques were offered in the study, and they were further classified. Security, key analysis, visual quality, and other measures were also used to evaluate existing VMEI approaches. The computational speed and visual quality of recreated images still have room for improvement, it was emphasized. Future research prospects and challenges in VMEI were discussed. In short, VMEI was still a developing area of study. Improving digital picture encryption might benefit from the findings of this state-of-the-art study. Wen, W.et al. (2016) in the present feature encryption techniques, edge information was considered crucial information. Images' salient portions often contain more crucial information than their edges. In order to create meaningful encrypted pictures, these authors research makes use of a salient regions encryption technique. To construct an encrypted image with visual significance, the author first identifies the salient parts using a saliency detection model in the compression domain, and then the author encrypts and decorates it. A visually





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relevant cipher text image cannot be decrypted to reveal the original image's prominent parts, as shown by experiments.

Y. Yang et al. (2023) the author suggested a lossless compression set partitioning in hierarchical trees (SPIHT) coding-based picture encryption system with visual significance. The first step was to jumble the wavelet coefficients of the original picture. The secret picture was then further compressed and scrambled using SPIHT coding. Finally, the cipher image was created by incorporating the concealed image into the cover art. Since a cipher picture does not seem like cipher text, the risk of an attack was lessened in a semi-honest trusted cloud environment. The experimental findings show that the suggested technique was fully reversible, which was an improvement over the prior algorithm. Yang, Y.-G. Et al. (2021) the author offers an M-ary decomposition and virtual bit VMIE technique. The best possible visual quality for VMEIs was these authors end aim. M-ary decomposition, M-ary corrected embedding, and the concepts of real and virtual bits were all essential to these authors methodology, which the author presented. The conventional, one-base representation of numbers was abandoned by these techniques. The bits in these authors algorithm were just conceptual. The author suggested a novel pre-encryption technique for images to accommodate these authors embedding model. The M-ary decomposition was employed in the pre-encryption process to handle plain pictures, and the resultant pre-encrypted images have a fixed pixel range. The non-standard pre-encryption technique was also found to have adequate security after an investigation was conducted.

DISCUSSION

The convergence of image encryption and digital signature technologies represents a watershed moment in addressing the critical needs of secure visual data transmission and storage. Unlike traditional encryption methods, which often result in distorted or incoherent image upon decryption, this ground-breaking methodology combines symmetric and asymmetric cryptographic algorithms, seamlessly interwoven with the power of digital signatures. This fusion not only strengthens data security but also rigorously maintains the visual content's integrity and coherence. This extensive review study methodically dissects the algorithm's underlying mathematics, exposing its complicated cryptographic foundations while clarifying its effectiveness in repelling adversary invasions. Beyond the theoretical, it highlights tangible applications in a variety of domains, ranging from healthcare, where patient data and diagnostic imagery can be securely handled without jeopardizing clinical acumen, to the media and entertainment industries, where copyrighted multimedia content is protected throughout the distribution chain. Furthermore, it is crucial in the sectors of government and military for preserving confidential visual data, including satellite images and intelligence briefings.

CONCLUSION

In a digital world where visual data security and meaning have become crucial, the combination of image encryption and digital signature technologies delivers a game-changing approach. Traditional image encryption technologies, although efficient in terms of data security, often rendered decrypted images aesthetically unreadable. In sharp contrast, the novel solution described in this survey study combines the strengths of cryptographic approaches, creating a symphony of symmetric and asymmetric encryption with the extra security layer of digital signatures. This study has shown the route toward attaining both strong security and visual coherence by thoroughly analyzing the algorithm's components, from its mathematical underpinnings through the encryption and decryption operations. Its practicality and importance are shown by its real-world applications in fields as varied as healthcare, multimedia, and national security.





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Table 1: comparison table for existing work

Author	Year	Methodology	Advantage	Limitation	
Armijo-	2020	Integer discrete	Unlike traditional image encryption	The modification of preprocessing and	
Correa,		wavelet	techniques that produce texture-like	embedding phases may introduce	
J.		transform	or noise-like encrypted images, this	increased computational complexity	
			approach generates visually	compared to traditional image	
			meaningful cover images.	encryption methods	
Chai, X.	2017	compressive	The technique was very secure	Compressive sensing algorithms can	
et al.		sensing	against both known-plaintext and	be computationally intensive,	
			chosen-plaintext assaults due to its	especially for large and high-	
			sensitivity to the plain picture.	resolution images.	
F. Jie et	2019	Integer	The pixel position permutation	Effective key management was critical	
al.		wavelet	based on the 2D-LASM sequence in	for the security of any encryption	
		transform	the first phase of the scheme adds	scheme.	
			an additional layer of security		
Hua, Z.	2021	Separable	The incorporation of a new parallel	The effectiveness of adaptive	
et al.		wavelet	CS technique markedly improves	techniques, such as adaptive-thres	
		transform	processing efficiency.	holding scarification, may depend on	
				selecting appropriate parameter	
				values.	
Jiang, D.	2021	Parallel	Optimizing data efficiency with the	Implementing the algorithm may	
et al.		compressive	use of parallel compressive sensing	require a deep understanding of the	
		sensing	(PCS) and the slant transform (ST).	underlying cryptographic and	
				mathematical concepts.	





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RESEARCH ARTICLE

Improving Deep Learning for Multiple Code Clone Detection through **Pretrained Models**

S Karthik and P Prabhakaran*

Assistant Professor, Department of Computer Science, PSG College of Arts & Science, (Affiliated to Bharathiar University), Coimbatore, Tamil Nadu, India.

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*Address for Correspondence

P Prabhakaran

Assistant Professor,

Department of Computer Science,

PSG College of Arts & Science, (Affiliated to Bharathiar University),

Coimbatore, Tamil Nadu, India.

Email: prabhakaranpsgcas@gmail.com



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ABSTRACT

Code-clone detection (CCD) is the technique of locating identical or similar chunks of code known as code clones within a program. Previously many machine learning (ML) techniques have been developed. These techniques always depend on inclusive, customized features to represent code fragments. So, handling multiple code clone (CC) labels will be a major problem that probably demands distinct evaluation criteria. In recent years, the research on CCD has been dramatically geared up by (DL) techniques. But, one most challenging issues faced by the DL approach in CCD is training from scratch for different domain source codes. Hence, in this research work, a transfer learning (TL) based attention learner (AL) model is utilized for the objective of code clone modelling (CCM). This model employs two pre-trained recurrent neural network (RNN)-based methods to transfer knowledge in the area of CCM rather than a single model. These pre-trained models are then implemented in TL as feature extractors. These collected characteristics are subsequently combined into an AL For application in various subsequent activities. Finally, AL applies the gained information of pre-trained models to fine-tune those features for specific down-streaming tasks. Finally, the experimental outcome demonstrates that the proposed TL-CCD technique has an accuracy of 96%, 98% and 97% for the dataset Apache Maven 3.8.3, Apache ant 1.10.12 and Opennlp-master 1.9.1, respectively, compared to the conventional CCM algorithms.

Keywords: Code-clone detection, Transfer learning, Attention learner, Recurrent Neural Network, Transfer knowledge





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INTRODUCTION

Code clones are code segments that are similar but originate in a different location. A significant amount of effort has gone into detecting, examining, and preventing code clones, which are ubiquitous in modern open-source software [1] and can have a range of negative implications. By considering the maintenance effort of clones, several methodologies and systems have been evolved to identify code clones. The methods are divided into groups based on the input, depiction, and algorithms as per the application of developers. These methods are broadly categorised as text-based [2, 3], token-based [4, 5], tree-based [6, 7], metric-based [8, 9], AST-based [10], Program Dependency Graph (PDG)-based [11], and hybrid methods [12]. While identifying all potential code clone pairings, these techniques function badly when dealing with complex source code architectures. To solve these issues for existing CCD techniques, the ML-based CCD has been developed. Generally, the ML process is very efficient and capable of detecting the code clone from the complex dataset using different models to reduce the software issues for the programmers in its preliminary stages itself. But still, the ML method was not suitable for detecting the multifeatured CC system. These multi-featured issues will be the major drawback of ML methods which possibly demands various evaluation criterion. For these types of issues, DL models play an important role in the efficient detection of CC. For example, instead of utilizing a predefined proximity metric or establishing a threshold for clone detection, the DL model is tailored to autonomously acquire the implicit code features from merged feature vectors to successfully identify clones among code snippets. But, the most difficult problem faced by the DL technique in CCD is training from scratch for different domain source codes. Motivated by this, this research devised a TL-AL-CCM technique that considerably enhances the efficiency of DL-based source code models. Initially, the concept of TL is employed in DL-based source code language models. The fundamental idea behind this research is to use a pre-trained source code language framework to transfer obtained information from one challenge to another. The two separate types of the RNN model are then trained for TL purposes. The learned information of pre-trained (RNN) models is integrated into an AL for a downstream activity. The AL applies pre-trained model learning information to given downstream tasks. In the TL model, pre-trained models are utilized to extract general features that may subsequently be fine-tuned for a specific purpose without having to train the model from scratch.

LITERATURE SURVEY

A weighted Recursive Auto-Encoder (RAE) was developed [13] for rapid CCD. Weighted RAE mined the program's attributes and encoded the functions to vectors before analyzing the program's abstract syntax tree. To improve the proportion of information donated by important nodes in the final vector representation of one application, the node weight information was taken into account in the abstract syntax tree during the modeling process. However, Threshold values greatly influence the performance of RAE. An effective detector called LV-Mapper was developed [14] to discover the LV-CCD using the third-generation sequencing alignment approach. Based on the locate-filterverify technique, this CCD may discover clones with more generic variance. The developed LV-Mapper locates and filters probable clone code pairings utilizing a restricted window of continuous lines known as seeds at a low cost. An adaptive point that fluctuated with program length was utilized for CCD. However, for LV-CCD, this method necessitates improvements in the refactoring and bug propagation systems. A unique technique of performing treebased convolution over a token-enhanced AST was proposed [15] to detect semantic clones. This method would explicitly gather sub-tree characteristics to fully exploit structural data in an AST. A token embedding method known as position-aware character embedding (PACE) was designed to boost the generalization ability for reducing the issue of unobserved data in CCD. However, this strategy resulted in a significant period and memory complexity. A ML approach was devised [16] for automatic CC validation. A training dataset is initially constructed by manually inspecting code clones collected from several clone detection techniques for distinct domain systems. Then, many attributes were retrieved to construct the ML model from those clones. The trained system was then utilized to identify clones without human intervention. Furthermore, the created model effectively reduces all false positive clones from the identification findings. However, this model performs poorly on smaller datasets. A novel





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approach of CCD was designed [17] that combines a deep feature learning system that trains the syntactic and semantic features with a code depiction created from the fusion embedding training of syntactic and semantic data at functional granularity to identify functional code clones. On the other hand, this technique performed less well on larger datasets. A functional code clone detector based on attention mechanism (FCCA) was introduced [18] for CCD. By maintaining both structured (code in the form of abstract syntax trees and control-flow graphs) and unstructured (code in the form of sequential tokens) data, this method was implemented on top of a cross-code representation. A collective representation was created by combining various code elements, and it included an attention mechanism that deliberately focused on key code components and attributes that improved the final detection accuracy. However, there were computational problems when this method was developed. A two-pass technique was developed [19] for CCD and its type categorization using a tree-based convolution neural network (TBCNN), which can capture structural features with a short propagation path and achieves comparable performance to word-by-word attention models. This technique detects clones on the first pass and then determines the clone type on the second pass. Both the first and second pass uses the TBCNN to extract features from the code fragments' ASTs to be compared. However, this method has high computational complexity issues.

A multi-threshold query, i.e. performing the query numerous times directed at distinct groups of clones, was implemented in a bag-of-tokens-based CCD [20] to find additional clone pairs of larger variation without reducing precision. To compensate for the additional behavior time generated by this technique, an optimization procedure was carried out to significantly minimize the duplication in detected clones. However, the classification accuracy of this approach was lower. An end-to-end cross-language (e2e-CL) CCD approach was developed [21] based on embeddings of ASTs using InferCode. In this model, InferCode captures the syntactical essence of a program as a vector. This model learns abstract features by grouping related embeddings through the contrastive loss function, which assures that comparable embeddings are close together but dissimilar embeddings are far apart. Moreover, the Siamese architecture was employed to overcome the data shortage and reduce overfitting issues. A new approach called Contrastive Cross-language model (CCL) was developed [22] for CCD. This model was efficient in detecting cross-language clones using learned representations. The pre-trained algorithm Code BERT was utilized in this approach to transform programs in many languages into high-dimensional vector representations. Furthermore, this system was improved using a provisional learning strategy that can discriminate between cloned and non-cloned pairs. However, this strategy produced minimal scalability.

PROPOSED WORK

This section discussed the brief illustration of TL-AL-CCM model. Figure 1 displays the complete workflow of the proposed model. Initially, the features like lexical, syntactic, semantic and structural features are extracted from the source code which is briefly provided in [23]. The implementation of the pre-trained algorithms for TL is then addressed. Once, the learned features are obtained, these features will be given as input to the AL. Furthermore, the AL method is provided to leverage and calibrate the obtained knowledge from pre-trained algorithms for the source code suggestion task. Next, this model is trained completely and it is tested by using the testing source code datsset.

Transfer Learning

One of the significant advancements in the DL paradigm for CCD is TL. By slightly adjusting the hyper-parameters, TL utilizes a pre-trained algorithm to conduct categorization on a dataset and then employs a similar strategy to another group of classified tasks. TL offers two advantages:

- 1. It can be learned faster because a model has already been learned on a distinct activity.
- 2. The model can be employed for tasks with smaller datasets because it has previously been learned on a larger dataset, and the weights are applied to the new tasks.

The TL technique for original code modelling is shown in Figure 2. The TL technique is utilized for the objective of effective source code modelling in code clones. This work proposes a distinct knowledge transfer approach, which transfers experience from two separate RNN-based pre-trained models and then fine-tunes it using an attention





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learner for a particular source code modelling activities. The proposed TL approach consists of different steps which are briefly listed below.

Pre-Training Models for Transfer Learning

In TL, pre-training is handled individually for each objective task, starting from scratch with each reference dataset which has been screened carefully. In actuality, pre-training will become more applicable over time in a variety of targeted activities, while starting from scratch for each one separately could reduce efficiency by allowing one to rediscover crucial data features. For the proposed task, two RNN-based models are pre-trained for TL in the CCM domain. Figure 3 depicts the structure of RNN model. In a given input sequence $U = (u_1,...,u_t)$, RNN upgrades its recurrent hidden state H_t by

$$H_t = \begin{cases} 0, & \text{if } T = 0\\ \varphi(H_{t-1}, u_t) & \text{otherwise} \end{cases}$$

$$\text{(1)}$$

$$W_t = \int_{0}^{\infty} \frac{1}{t^2} \left(\frac{1}{t^2} + \frac{1}{t^2} \frac{1}{t^2} \right) dt = \int_{0}^{\infty} \frac{1}{t^2} \left(\frac{1}{t^2} + \frac{1}{t^2} \frac{1}{t^2} \right) dt = \int_{0}^{\infty} \frac{1}{t^2} dt = \int_{0}^{\infty} \frac{1}{$$

 u_t and H_t denote the data input and recurrent hidden state, with time step t, respectively, $\phi(.)$ denotes the non-linear activation function of a hidden layer, such as sigmoid or hyperbolic tangent. The following is the updated rule for the recurrent concealed state in (1):

$$H_t = \varphi \left(Z_{u_t} + A(H_{t-1}) \right) \tag{2}$$

Where Z and A are the input co-efficient vectors at the present stage, and the activation of the H_t is provided in the previous step, respectively. Equation (2) is expanded as follows:

$$H_t = \varphi(Z_{pHu_t} + Z_{HH}H_{t-1} + B_H) \tag{3}$$

Where, t = 1 to T, Z_{PH} represents an input-hidden weight vector, Z_{HH} and B_H represents the weight matrix of the hidden layer, and hidden layer bias vector respectively. However, it has been found that one significant problem in RNNs is the vanishing gradient, which can be rectified by employing the gated recurrent unit. The GRU avoids the vanishing gradient problem by revealing the complete hidden data on each time step t. It can be said to be

$$H_p = (1 - w_p)H_{p-1} + w_pH_p \tag{4}$$

$$w_p = \phi(Z_w T_p + A_w(H_{p-1})) \tag{5}$$

$$H_p = \tanh(ZT_p + v_p \otimes A H_{p-1}) \tag{6}$$

$$v_p = \phi(Z_v T_p + A_v H_{p-1}) \tag{7}$$

w and v are specified as the reset gate and the update gate, respectively. The gate operations like p will restrain the inflating of the memory gradient and reproduce the gradient from a linear multiplication expression to a summation expression. This helps to resolve the problems, particularly, the vanishing gradient.

Training the model to transfer knowledge

For TL, the pre-trained RNN model is described in the earlier section. Then, the learned data knowledge is converted using these pre-trained models for the subsequent function of source code suggestion. The major objective is to maintain the gathered knowledge in pre-trained models, freeze the extracted knowledge, and then fine-tune it for the source code suggestion activity. To attain maximum performance, the attention learner focuses on task-particular extracted aspects (lexica, syntactic, semantic, and structural). The objective of an AL process is to advise the model during training on which portion of the input data is crucial so that it can pay close attention to the data. The pre-trained RNN layer can process and generate the learned features from the source code, the length of the learned feature of hidden state vectors is always equal to the length of the input (extracted) features. The alignment among two sequences is constructed to train the model for the distribution of s ($j_1, j_2, ..., j_l$ | $H_1, H_2, ..., H_p$) for which there is no clear functional dependency between I and P. The RNN-based AL model produces the resulted features of ($j_1, j_2, ..., j_l$) one time at a time, simultaneously aligning each generated element to the encoded input feature ($H_1, H_2, ..., H_p$). The composition of RNN and AL works efficiently for modelling the clone codes. The AL selects the temporal positions over the learned features which should be used to update the hidden state of the RNN and to make a prediction for the next output value. Typically, the selection of elements from the input features is a weighted sum

$$E_t = \sum_M A_{tM} H_M \tag{8}$$

Where, A_tM is the attention weights and it should satisfy $A_{tM}>0$ and $\sum_{M} A_{tM}=1$





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In this process, The AL process is used for tine-tuning the learned features from pre-trained RNN layer for modelling the code clones. The determination is defined in (9), (10) and (11)

$$=R^*A_{t-1} \tag{9}$$

$$f_{tM} = z^T \tanh (X_{l_{t-1}} + YH_M + W_{g_M} + C)$$
 (10)

$$A_{tM} = e(f_{tM}) / \sum_{M=1}^{m} e(f_{tM})$$
(11)

Where, X, Y, W and R are the feature matrix, z and C are the feature vector, * denotes convolution $l_{(l-1)}$ stands for the former RNN component of the AL mechanism. The equation (11) depicts the weight A_{LM} obtained by normalizing the scores f_{LM} . In Equation (3), the resulting score depends on the previous state $l_{(l-1)}$, the feature label in the respective position H_M and the vector of so-called convolutional features g_M . The name "convolutional" comes from the convolution along the time axis used in equation (2) to compute the matrix G that comprises all feature vectors g_M . Similarly, the AL mechanism describes above combines source code information from $l_{(l-1)}$, the feature in the respective position H_M and focuses on the previous step described by attention weight $A_{(l-1)}$, which particularly defines A_l is important for code modelling.

Fine-tuning the learned model for code clone modelling

The figure 4 completely describes the fine-tuning process of learned features in TL model with AL mechanism for CCM. In figure 4, after the completion of transferring trained features from the pre-trained RNN model, the features obtained from the AL mechanism are considered to train the TL-CCD model and the trained model is employed to predict the code clones from the testing dataset which is taken from the source code. Fine-tuning is the process of unfreezing different stages in the pre-trained model (learned features). The TL is typically carried out in conjunction with the DL model to effectively calibrate the pre-trained model on the original code using certain data from the target task. By using this concept in this research, the learned features from the pre-trained models are fine-tuned to model the code clones.

RESULT AND DISCUSSION

assess the effectiveness of the suggested method and to detect and model the clone and its type using its similarity attributes. A benchmark dataset, such as Big Clone Bench, is used to generate the training dataset. The Big Clone Bench is a benchmark consisting of a large number of manually approved clones from the IJaDataset-2.0 source. The testing dataset is then gathered from Apache Maven 3.8.3 is a toolkit for managing and comprehending software projects. Maven depends upon the presumption of a Project Entity Paradigm (POM), and can manage a project's development, monitoring, and information from a primary source of contact [24]. Apache ant 1.10.12 is a Java library and function tool that manages operations indicated in construct documents as objectives and extensibility endpoints. The well-known use of Ant is the creation of Java applications, as it has a collection of built-in tasks for compiling, assembling, testing, and executing Java programmes [25]. Apache Opennlp-master 1.9.1 is a natural language information retrieval toolkit based on ML techniques. Among the most common NLP operations supported are indexing, sentence classification, part-of-speech labeling, named entity identification, stacking, and processing [26] On using these datasets, the efficiency of proposed TL-AL-CCM technique is compared with an existing method TBCNN-CCD [19], e2e-CL- CCD [21] and CCL-CCD [22] in terms of accuracy, precision, recall, time period, Memory space and Clone types for detecting and modelling all four types of CC with its similarity features.

For the experimental purpose, the dataset is divided into two different categories as a training and testing dataset to

Accuracy

It is calculated by dividing the number of properly-recognized CCs by the absolute No. of CCs and non-CCs in the cloned repository (actual). Table 1 provides a comparison of existing and proposed methods for detecting and modelling all clone types. Figure 5 displays the results of accuracy achieved for proposed and existing methods to improve the CCM task. From the above analysis, it is observed that the accuracy of the proposed model is 11.28%, 8.04%, and 3.53% higher than the TBCNN-CCD, e2e-CL- CCD and CCL-CCD for Apache Maven 3.8.3 dataset;





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15.27%, 6.74%, and 4.03% higher than the TBCNN-CCD, e2e-CL- CCD and CCL-CCD for Apache ant 1.10.12 dataset; 13.70%, 7.77%, and 2.32% higher than the TBCNN-CCD, e2e-CL- CCD and CCL-CCD for Opennlp-master datasets respectively. From this analysis, it is proved that the proposed model attains higher accuracy than the other existing methods for efficient CCM tasks.

Precision

It is computed by calculating the overall No. of identified CCs by the No. of accurately detected CCs (predicted). Table 2 compares extant and proposed methods for identifying and modifying all clone variants. Figure 6 displays the results of precision achieved for proposed and existing methods to improve the CCM tasks. From the above analysis, it is observed that the precision of the proposed model is 18.02%, 13.88%, and 6.78% higher than the TBCNN-CCD, e2e-CL- CCD and CCL-CCD for Apache Maven 3.8.3 dataset; 20.15%, 14.02%, and 6.14% higher than the TBCNN-CCD, e2e-CL- CCD and CCL-CCD for Apache ant 1.10.12 dataset; 17.69%, 13.04%, and 4.75% higher than the TBCNN-CCD, e2e-CL- CCD and CCL-CCD for Opennlp-master datasets respectively. From this analysis, it is proved that the proposed model attains a higher precision than the other existing methods for the efficient CCM tasks

Recall

The recall is obtained by splitting the total count of CCs in the database by the number of successfully recognized CCs (actual). Table 3 provides a comparison of existing and proposed methods for detecting all clone types. Figure 7 displays the results of recall achieved for proposed and existing methods to improve the CCM tasks. From the above analysis, it is observed that the recall of proposed model is 13.96%, 8.06%, and 4.48% higher than the TBCNN-CCD, e2e-CL- CCD and CCL-CCD for Apache Maven 3.8.3 dataset; 16.51%, 9.89%, and 6.93% higher than the TBCNN-CCD, e2e-CL- CCD and CCL-CCD for Apache ant 1.10.12 dataset; 13.02%, 7.57%, and 4.25% higher than the TBCNN-CCD, e2e-CL- CCD and CCL-CCD for Opennlp-master datasets respectively. From this analysis, it is proved that the proposed model attains a higher recall than the other existing methods for efficient CCM tasks.

CONCLUSION

In this research work, TL-AL-CCM technique enhances the performance of DL-based source code models greatly. Initially, the principle of TL is applied in DL-based source code language models. Following that, the two RNN-based pre-trained models are used for TL in the region of reference code. The pre-trained models' learned information is integrated into an AL for subsequent issues. The AL applies pre-trained model learning data to given downstream activities. In TL, pre-trained models are employed to capture generic attributes that can subsequently be calibrated for a specific purpose without having to retrain the model from scratch. At last, the suggested method provides better results than other existing methods in terms of accuracy, precision, and recall for three different datasets Apache Maven 3.8.3 Apache ant 1.10.12 dataset Opennlp-master.

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Table 1 Evaluation of Accuracy

Datasets	TBCNN-CCD	e2e-CL- CCD	CCL-CCD	TL-AL-CCM
Apache Maven 3.8.3	0.869	0.895	0.934	0.967
Apache ant 1.10.12	0.851	0.919	0.943	0.981
Opennlp-master 1.9.1	0.854	0.901	0.949	0.973





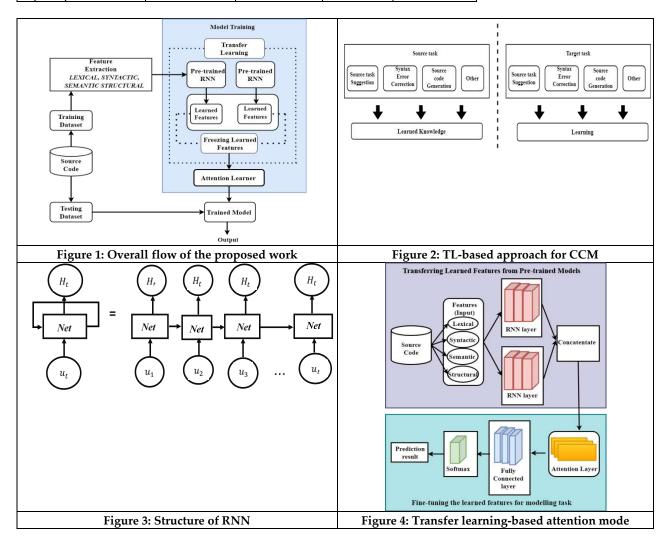
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Table 2 Evaluation of Precision

Datasets	TBCNN-CCD	e2e-CL- CCD	CCL-CCD	TL-AL-CCM
Apache Maven 3.8.3	0.827	0.857	0.914	0.976
Apache ant 1.10.12	0.819	0.863	0.927	0.984
Opennlp-master 1.9.1	0.825	0.859	0.927	0.971

Table 3 Evaluation of Recall

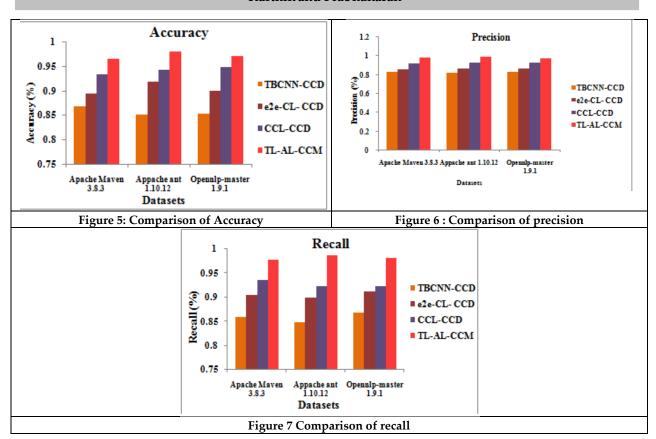
Datasets	TBCNN-CCD	e2e-CL- CCD	CCL-CCD	TL-AL-CCM
Apache Maven 3.8.3	0.859	0.906	0.937	0.979
Appache ant 1.10.12	0.848	0.899	0.924	0.988
Opennlp-master 1.9.1	0.868	0.912	0.941	0.981







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RESEARCH ARTICLE

Mathematical Analysis of New SEIQR Model via Laplace Adomian **Decomposition Method**

N. Jeeva^{1*} and K.M. Dharmalingam²

¹Research Scholar, Department of Mathematics, The Madura College (Affiliated to Madurai Kamaraj University) Madurai, Tamil Nadu, India.

²Associate Professor, Department of Mathematics, The Madura College (Affiliated to Madurai Kamaraj University) Madurai, Tamil Nadu, India.

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*Address for Correspondence

N. Jeeva

Research Scholar,

Department of Mathematics,

The Madura College (Affiliated to Madurai Kamaraj University)

Madurai, Tamil Nadu, India.

Email: jeevapirc2405@gmail.com



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ABSTRACT

This study focuses on a novel mathematical model for the coronavirus, building upon the SEIQR epidemiological framework while incorporating a unique compartment for isolation or quarantine. We employed the Laplace Adomian decomposition Method (LADM) to perform an in-depth analytical examination of this novel model. The effectiveness of the LADM was demonstrated by providing highly accurate approximate analytical solutions with minimal iterations. In addition, we conducted a comparative analysis using numerical simulations to comprehensively understand the behaviour of the model. This comparative approach not only validates the results but also provides valuable insights into how the novel coronavirus model behaves under various conditions and scenarios.

Keywords: Novel Corona Virus transmission, SEIQR model, Laplace Adomian Decomposition Method (LADM), Mathematical modelling, Numerical simulation.

INTRODUCTION

The term "coronavirus" denotes a family of viruses capable of causing illness in both animals and humans. The novel coronavirus, formally designated as "SARS-CoV-2" (Severe Acute Respiratory Syndrome Coronavirus 2), was initially detected in Wuhan, China, in December 2019. It is accountable for the onset of "COVID-19" (Coronavirus Disease 2019) [1]. This virus has left a profound impact on public health and has triggered a worldwide pandemic.





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Measures to curb its spread encompass vaccination campaigns, adherence to public health protocols such as maskwearing and social distancing, and ongoing research to enhance our comprehension of the virus and its various strains [2-3]. The SIR model is a well-established mathematical framework in epidemiology, extensively employed for analyzing the transmission of infectious diseases within a population. The acronym "SIR" denotes the three compartments used to classify individuals within the population: Susceptible, Infected, and Recovered [4-5]. SEIQR serves as an epidemiological framework employed to investigate the propagation of infectious diseases within a population. It represents an expansion of the fundamental SIR (Susceptible-Infectious-Recovered) model, introducing supplementary compartments to accommodate diverse stages of disease transmission. The SEIQR model conventionally encompasses the following compartments: Susceptible, Exposed, Infectious, Quarantined, and Recovered [6-8]. This model proves especially valuable when examining diseases characterized by a substantial incubation period, during which infected individuals are not yet capable of transmitting the infection.

Its utility lies in its capacity for a more intricate examination of disease transmission dynamics, offering insights into forecasting the future course of an outbreak and assessing the impact of interventions, such as quarantine measures and vaccination campaigns [9]. The mathematical model for the transmission of the novel coronavirus was formulated in a study conducted by researchers [10]. This mathematical model employs a collection of non-linear differential equations to incorporate the propagation of COVID-19 infection. It includes a unique compartment class for isolation or quarantine and estimates the model parameters by aligning the model with reported data on the evolving pandemic situation in India. The primary objective of this paper is to utilize analytical method to solve the model for the novel coronavirus. The approximate analytical solutions for the novel coronavirus model are obtained using a potent technique known as the Laplace Adomian Decomposition Method (LADM). This method yields highly precise results with minimal iterations, making it a valuable tool for analyzing the dynamics of the novel coronavirus model. The LADM involves breaking down the equations into infinite power series of Adomian polynomials, simplifying the computation of approximate solutions for the model's variables.

Mathematical Modelling of the problem

In [10], a deterministic novel model for Corona Virus was introduced, utilizing a system of ordinary non-linear differential equations. The population was divided into five compartments: Susceptible population S(t), Exposed population E(t), Infected population I(t), Isolated population Q(t), and Recovered population R(t), as illustrated in Figure 1. The Novel SEIQR model for Corona Virus can be expressed as follows:

$$\frac{dS}{dt} = A - \beta SI - \mu S \tag{1}$$

$$\frac{dt}{dt} = A - \beta SI - \mu S \tag{1}$$

$$\frac{dE}{dt} = \beta SI - (\alpha + \mu)E \tag{2}$$

$$\frac{dI}{dt} = \alpha E - (\gamma + \alpha + \mu)I \tag{3}$$

$$\frac{\tilde{d}\tilde{l}}{dt} = \alpha E - (\gamma + q + \mu)I \tag{3}$$

$$\frac{dQ}{dQ} = qI - (\theta + \mu)Q \tag{4}$$

$$\frac{dl}{dt} = \alpha E - (\gamma + q + \mu)I$$
(3)
$$\frac{dQ}{dt} = qI - (\theta + \mu)Q$$
(4)
$$\frac{dR}{dt} = \gamma I + \theta Q - \mu R$$
(5)

Subject to the initial conditions:

$$S(0) = a, E(0) = b, I(0) = c, Q(0) = d, R(0) = e$$

$$(6)$$

Where, Table 1 provides a list of the parameter representations utilized in this SEIQR model.

Approximate Analytical solution using LADM

The Laplace Adomian Decomposition Method (LADM) is a mathematical method employed to address differential equations, particularly those characterized by nonlinearity. LADM integrates the fundamental concepts of Laplace transforms and the Adomian Decomposition Method (ADM) to establish a structured framework for deriving analytical solutions to a diverse array of differential equations. An important and noteworthy advantage of LADM is its ability to address nonlinear differential equations without the need for linearization, perturbation, transformation, or any form of discrimination. With LADM, it yields an analytical solution in the form of a rapidly converging infinite power series. Its effectiveness extends to various domains, encompassing fluid dynamics, heat transfer,





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population dynamics, and more. Across these fields, LADM has consistently proven its capability to effectively handle complex problems and offer valuable insights [11-17].

Basic concept of Laplace Adomian Decomposition Method (LADM)

In this sub-section, we explore the application of a hybrid method that combines the Laplace transformation and the Adomian decomposition algorithm to solve the nonlinear autonomous first-order differential equation that governing the problem.

$$L[u(x)] + R[u(x)] + N[u(x)] = g(x)$$
(7)

$$u(0) = f(x) \tag{8}$$

$$L[u(x)] = g(x) - R[u(x)] - N[u(x)]$$
(9)

By applying the Laplace transform to both sides of Eq. (9) and utilizing the differentiation property, we obtain the following result.

$$sL[u(x)] - f(x) = L[g(x)] - L[R[u(x)]] - L[N[u(x)]]$$

$$L[u(x)] = \frac{1}{s}f(x) + \frac{1}{s}L[g(x)] - \frac{1}{s}L[R[u(x)]] - \frac{1}{s}L[N[u(x)]]$$
(10)

When we apply the inverse Laplace transform to both sides of Eq. (10), we obtain the following result.

$$u(x) = \varphi(x) - L^{-1} \left[\frac{1}{s} L[R[u(x)]] - \frac{1}{s} L[N[u(x)]] \right]$$
(11)

Here, $\varphi(x)$ represents the term originating from the first three terms on the right-hand side of Eq. (11). Next, we make an assumption about the solution to the problem in the form of a decomposing series.

$$u(x) = \sum_{n=0}^{\infty} u_n(x) \tag{12}$$

Likewise, the nonlinear terms are expressed in relation to the Adomian polynomials as follows:

$$N[u(x)] = \sum_{n=0}^{\infty} A_n \tag{13}$$

Where the A_n 's represents the Adomain polynomials defined in the form

$$A_n = \frac{1}{n!} \frac{d^n}{d\lambda^n} \left[N(\sum_{n=0}^{\infty} \lambda^n y_n) \right] |_{\lambda=0}$$

$$\tag{14}$$

Plugging Eqn.(12) and (13) into Eq.(14), we obtain

$$\sum_{n=0}^{\infty} u_n(x) = \varphi(x) - L^{-1} \left[\frac{1}{\varsigma} L[R \sum_{n=0}^{\infty} u_n(x)] - \frac{1}{\varsigma} L[N \sum_{n=0}^{\infty} A_n] \right]$$
(15)

By equating both sides of Eq.(15), we derive an iterative algorithm in the following form:

$$u_0(x)=\varphi(x)$$

$$\begin{split} u_1(x) &= -L^{-1} \big[\frac{1}{s} L \left[R \sum_{n=0}^{\infty} u_0(x) \right] - \frac{1}{s} L \left[N \sum_{n=0}^{\infty} A_0 \right] \big] \\ u_2(x) &= -L^{-1} \big[\frac{1}{s} L \left[R \sum_{n=0}^{\infty} u_1(x) \right] - \frac{1}{s} L \left[N \sum_{n=0}^{\infty} A_1 \right] \big] \end{split}$$

$$u_{n+1}(x) = -L^{-1}\left[\frac{1}{\varsigma}L[R\sum_{n=0}^{\infty}u_n(x)] - \frac{1}{\varsigma}L[N\sum_{n=0}^{\infty}A_n]\right]$$
(16)

The solution to the differential equation is then obtained as the summation of the decomposed series in the following form:

$$u(x) \approx u_0(x) + u_1(x) + u_2(x) + \cdots$$
 (17)

Approximate Analytical solution of Covid-19 SEIQR model using LADM

In this sub-section, We illustrate the application of the Laplace-Adomian Decomposition Method to nonlinear ordinary differential systems. To initiate the process, we commence by applying the Laplace transformation, denoted by L, to both sides of equations (1-5).

$$L\left[\frac{dS}{dt}\right] = L[A] - L[\beta SI] - L[\mu S] \tag{18}$$

$$L\left[\frac{dS}{dt}\right] = L[A] - L[\beta SI] - L[\mu S]$$

$$L\left[\frac{dE}{dt}\right] = L[\beta SI] - L[(\alpha + \mu)E]$$
(18)

$$L\left[\frac{dI}{dt}\right] = L[\alpha E] - L[(\gamma + q + \mu)I] \tag{20}$$

$$L\left[\frac{dQ}{dL}\right] = L[qI] - L[(\theta + \mu)Q] \tag{21}$$

$$L\left[\frac{dR}{dt}\right] = L[\gamma I] + L[\theta Q] - L[\mu R] \tag{22}$$

We make use of the properties of derivatives and constants in the Laplace transform for our analysis.





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$$wL[S] - S(0) = \frac{A}{w} - \beta L[SI] - \mu L[S]$$
(23)

$$wL[E] - E(0) = \beta L[SI] - (\alpha + \mu)L[E]$$
(24)

$$wL[I] - I(0) = \alpha L[E] - (\gamma + q + \mu)L[I]$$
(25)

$$wL[Q] - Q(0) = qL[I] - (\theta + \mu)L[Q]$$
(26)

$$wL[R] - R(0) = \gamma L[I] + \theta L[Q] - \mu L[R]$$
(27)

Upon applying the initial conditions, we obtain the following results.

$$L[S] = \frac{a}{w} + \frac{A}{w^2} - \frac{\beta}{w} L[SI] - \frac{\mu}{w} L[S]$$
 (28)

$$L[E] = \frac{b}{w} + \frac{\beta}{w} L[SI] - \frac{(\alpha + \mu)}{w} L[E]$$
(29)

$$L[I] = \frac{c}{\Box} + \frac{\alpha}{\Box} L[E] - \frac{(\gamma + q + \mu)}{\Box} L[I] \tag{30}$$

$$L[S] = \frac{a}{w} + \frac{A}{w^2} - \frac{\beta}{w} L[SI] - \frac{\mu}{w} L[S]$$
(28)
$$L[E] = \frac{b}{w} + \frac{\beta}{w} L[SI] - \frac{(\alpha + \mu)}{w} L[E]$$
(29)
$$L[I] = \frac{c}{w} + \frac{\alpha}{w} L[E] - \frac{(\gamma + q + \mu)}{w} L[I]$$
(30)
$$L[Q] = \frac{d}{w} + \frac{q}{w} L[I] - \frac{(\theta + \mu)}{w} L[Q]$$
(31)
$$L[R] = \frac{e}{w} + \frac{\gamma}{w} L[I] + \frac{\theta}{w} L[Q] - \frac{\mu}{w} L[R]$$
(32)

$$L[R] = \frac{e}{w} + \frac{\gamma}{w} L[I] + \frac{\theta}{w} L[Q] - \frac{\mu}{w} L[R]$$
(32)

We make the assumption that the solutions for S, E, I, Q, and R can be expressed as infinite series. Additionally, we decompose the nonlinear terms involving SI = U in the model using Adomian polynomials, which are expressed as follows:

$$S = \sum_{n=0}^{\infty} S_n, \ E = \sum_{n=0}^{\infty} E_n, \ I = \sum_{n=0}^{\infty} I_n, \ Q = \sum_{n=0}^{\infty} Q_n, R = \sum_{n=0}^{\infty} R_n, U = \sum_{n=0}^{\infty} U_n$$
(33)

The Adomian polynomials, denoted by A_n , are defined as follows;

$$U_n = \frac{1}{n!} \frac{d^n}{d\lambda^n} \left[\sum_{i=0}^{\infty} \lambda^i S_i \sum_{i=0}^{\infty} \lambda^i I_i \right]$$

$$U_0 = S_0 I_0$$

$$U_1 = S_0 I_1 + S_1 I_0$$
(34)

Upon inserting eqn.4 and eqn.5 into eqn.3, we obtain the following:

$$L[\sum_{n=0}^{\infty} S_n] = \frac{a}{w} + \frac{A}{w^2} - \frac{\beta}{w} L[\sum_{n=0}^{\infty} U_n] - \frac{\mu}{w} L[\sum_{n=0}^{\infty} S_n]$$
(35)

Upon inserting eqn.4 and eqn.5 into eqn.5, we obtain the following:
$$L[\sum_{n=0}^{\infty} S_n] = \frac{a}{w} + \frac{A}{w^2} - \frac{\beta}{w} L[\sum_{n=0}^{\infty} U_n] - \frac{\mu}{w} L[\sum_{n=0}^{\infty} S_n]$$
(35)

$$L[\sum_{n=0}^{\infty} E_n] = \frac{b}{w} + \frac{\beta}{w} L[\sum_{n=0}^{\infty} U_n] - \frac{(\alpha + \mu)}{w} L[\sum_{n=0}^{\infty} E_n]$$
(36)

$$L[\sum_{n=0}^{\infty} I_n] = \frac{c}{w} + \frac{a}{w} L[\sum_{n=0}^{\infty} E_n] - \frac{(\gamma + q + \mu)}{w} L[\sum_{n=0}^{\infty} I_n]$$
(37)

$$L[\sum_{n=0}^{\infty} Q_n] = \frac{d}{w} + \frac{q}{w} L[\sum_{n=0}^{\infty} I_n] - \frac{(\theta + \mu)}{w} L[\sum_{n=0}^{\infty} Q_n]$$
(38)

$$L[\sum_{n=0}^{\infty} R_n] = \frac{e}{w} + \frac{\gamma}{w} L[\sum_{n=0}^{\infty} I_n] + \frac{\theta}{w} L[\sum_{n=0}^{\infty} Q_n] - \frac{\mu}{w} L[\sum_{n=0}^{\infty} R_n]$$
(39)

$$L[\sum_{n=0}^{\infty} I_n] = \frac{c}{w} + \frac{\alpha}{w} L[\sum_{n=0}^{\infty} E_n] - \frac{(\gamma + q + \mu)}{w} L[\sum_{n=0}^{\infty} I_n]$$
(37)

$$L[\sum_{n=0}^{\infty} Q_n] = \frac{d}{w} + \frac{q}{w} L[\sum_{n=0}^{\infty} I_n] - \frac{(\theta + \mu)}{w} L[\sum_{n=0}^{\infty} Q_n]$$
(38)

$$L[\sum_{n=0}^{\infty} R_n] = \frac{e}{w} + \frac{\gamma}{w} L[\sum_{n=0}^{\infty} I_n] + \frac{\theta}{w} L[\sum_{n=0}^{\infty} Q_n] - \frac{\mu}{w} L[\sum_{n=0}^{\infty} R_n]$$
(39)

The initial iteration of the Laplace-Adomian Decomposition Method is obtained as follows:

$$L[S_0] = \frac{a}{w} + \frac{A}{w^2} \tag{40}$$

$$L[S_0] = \frac{a}{w} + \frac{A}{w^2}$$

$$L[E_0] = \frac{b}{w}$$
(40)

$$L[I_0] = \frac{\ddot{c}}{w} \tag{42}$$

$$L[Q_0] = \frac{d}{w} \tag{43}$$

$$L[Q_0] = \frac{d}{w}$$

$$L[R_0] = \frac{e}{w}$$

$$(43)$$

Applying the inverse Laplace transform to the initial iteration yields the final solution.

$$S_0 = a + At (45)$$

$$E_0 = b \tag{46}$$

$$I_0 = c (47)$$

$$Q_0 = d \tag{48}$$

$$R_0 = e (49)$$





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The first iteration of the Laplace-Adomian Decomposition Method is derived by comparing both sides of the equation.

$$L[S_1] = -\frac{\beta}{w} L[U_0] - \frac{\mu}{w} L[S_0] \tag{50}$$

$$L[E_1] = \frac{\beta}{w} L[U_0] - \frac{(\alpha + \mu)}{w} L[E_0]$$
 (51)

$$L[I_1] = -\frac{\alpha}{w} L[E_0] - \frac{(\gamma + q + \mu)}{w} L[I_0]$$
(52)

$$L[S_{1}] = -\frac{\beta}{w} L[U_{0}] - \frac{\mu}{w} L[S_{0}]$$

$$L[E_{1}] = \frac{\beta}{w} L[U_{0}] - \frac{(\alpha + \mu)}{w} L[E_{0}]$$

$$L[I_{1}] = \frac{\alpha}{w} L[E_{0}] - \frac{(\gamma + q + \mu)}{w} L[I_{0}]$$

$$L[Q_{1}] = \frac{q}{w} L[I_{0}] - \frac{(\theta + \mu)}{w} L[Q_{0}]$$

$$L[R_{1}] = \frac{\gamma}{w} L[I_{0}] + \frac{\theta}{w} L[Q_{0}] - \frac{\mu}{w} L[R_{0}]$$
(54)

$$L[R_1] = \frac{\ddot{\gamma}}{L}[I_0] + \frac{\theta}{L}[Q_0] - \frac{\mu}{L}[R_0] \tag{54}$$

By utilizing the initial iteration, we obtain the following.

$$L[S_1] = -\frac{\mu a}{w^2} - \frac{[\beta ac + A\mu]}{w^3} - \frac{\beta Ac}{w^4}$$
 (55)

by utilizing the initial iteration, we obtain the following.

$$L[S_1] = -\frac{\mu a}{w^2} - \frac{[\beta ac + A\mu]}{w^3} - \frac{\beta Ac}{w^4}$$
(55)

$$L[E_1] = -\frac{(\alpha + \mu)b}{w^2} + \frac{\beta ac}{w^3} + \frac{\beta Ac}{w^4}$$
(56)

$$L[I_1] = \frac{ab - (\gamma + q + \mu)c}{w^2}$$
(57)

$$L[I_1] = \frac{ab - (\gamma + q + \mu)c}{w^2} \tag{57}$$

$$L[Q_1] = \frac{qc - (\ddot{\theta} + \mu)d}{m^2} \tag{58}$$

$$L[I_1] = \frac{w^2}{w^2}$$

$$L[Q_1] = \frac{qc - (\theta + \mu)d}{w^2}$$

$$L[R_1] = \frac{\gamma c + \theta d - \mu e}{w^2}$$
(59)

Similarly, for the first iteration.

$$S_{1} = -\mu at - \left[\beta ac + A\mu\right] \frac{t^{2}}{2!} - \beta Ac \frac{t^{3}}{3!}$$

$$E_{1} = -(\alpha + \mu)bt + \beta ac \frac{t^{2}}{2!} + \beta Ac \frac{t^{3}}{3!}$$

$$(60)$$

$$E_1 = -(\alpha + \mu)bt + \beta ac \frac{t^2}{2!} + \beta Ac \frac{t^3}{3!}$$
(61)

$$I_1 = (\alpha b - (\gamma + q + \mu)c)t \tag{62}$$

$$Q_1 = (qc - (\theta + \mu)d)t \tag{63}$$

$$R_1 = (\gamma c + \theta d - \mu e)t \tag{64}$$

In a similar way, the second iteration of the Laplace-Adomian Decomposition Method is derived by comparing both sides of the equation.

sides of the equation:

$$L[S_2] = -\frac{\beta}{w} L[U_1] - \frac{\mu}{w} L[S_1]$$
(65)

$$L[E_2] = \frac{\beta}{w} L[U_1] - \frac{(\alpha + \mu)}{w} L[E_1]$$
(66)

$$L[I_2] = \frac{\alpha}{w} L[E_1] - \frac{(\gamma + q + \mu)}{w} L[I_1]$$
(67)

$$L[Q_2] = \frac{q}{w} L[I_1] - \frac{(\theta + \mu)}{w} L[Q_1]$$
(68)

$$L[R_2] = \frac{\gamma}{w} L[I_1] + \frac{\theta}{w} L[Q_1] - \frac{\mu}{w} L[R_1]$$
(69)

$$L[E_2] = \frac{\beta}{w} L[U_1] - \frac{(\alpha + \mu)}{w} L[E_1]$$
(66)

$$L[I_2] = \frac{\alpha}{w} L[E_1] - \frac{(\gamma + q + \mu)}{w} L[I_1]$$
(67)

$$L[Q_2] = \frac{q}{w} L[I_1] - \frac{(\theta + \mu)}{w} L[Q_1]$$
(68)

$$L[R_2] = -\frac{\eta}{w} L[I_1] + \frac{\theta}{w} L[Q_1] - \frac{\mu}{w} L[R_1]$$
(69)

By utilizing the initial iteration and first iteration, we obtain the following.

$$S_{2} = \mu^{2} a \frac{t^{2}}{2!} + \left(\beta \mu a c - \beta a \left(\alpha b - (\gamma + q + \mu)\right) + \mu (\beta a c + A \mu)\right) \frac{t^{3}}{3!} + \left(\beta c (\beta a c + A \mu) + \beta A \left(\alpha b - (\gamma + q + \mu)\right) + \beta \mu A c\right) \frac{t^{4}}{4!} + \beta^{2} A c^{2} \frac{t^{5}}{5!}$$
(70)

$$E_{2} = (\alpha + \mu)^{2} b \frac{t^{2}}{2!} + \left(-(\alpha + \mu)\beta ac - \beta \mu ac + \beta a \left(\alpha b - (\gamma + q + \mu)\right)\right) \frac{t^{3}}{3!} + \left(\beta A \left(\alpha b - (\gamma + q + \mu)\right) - \beta c (\beta ac + A\mu) - \alpha + \mu \beta ac t 44! - \beta 2 Ac 2t 55!\right)$$
(71)

$$I_{2} = \left(-\alpha(\alpha + \mu)b + (\gamma + q + \mu)(\alpha b - (\gamma + q + \mu)c)\right) \frac{t^{2}}{2!} + \alpha \beta a c \frac{t^{3}}{3!} + \alpha \beta A c \frac{t^{4}}{4!}$$

$$Q_{2} = (q(\alpha b - (\gamma + q + \mu)c) - (\theta + \mu)(qc - (\theta + \mu)d)) \frac{t^{2}}{2!}$$
(72)

$$Q_2 = (q(\alpha b - (\gamma + q + \mu)c) - (\theta + \mu)(qc - (\theta + \mu)d))\frac{t^2}{2!}$$
(73)

$$R_2 = (\gamma(\alpha b - (\gamma + q + \mu)c) + \theta(qc - (\theta + \mu)d) - \mu(\gamma c + \theta d - \mu e))\frac{t^2}{2!}$$
(74)

Subsequently, the remaining terms can be computed using a similar approach. By utilizing these calculated values, we can approximate the solutions to the above systems in the form of an infinite series as follows.

$$S(t) = S_0 + S_1 + S_2 + \cdots (75)$$



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$$E(t) = E_0 + E_1 + E_2 + \cdots$$

$$I(t) = I_0 + I_1 + I_2 + \cdots$$

$$Q(t) = Q_0 + Q_1 + Q_2 + \cdots$$

$$R(t) = R_0 + R_1 + R_2 + \cdots$$
(76)
(77)

Therefore, the approximate analytical solutions of the SEIQR equations are derived by substituting Eqns.(45-49), Eqn.(60-64) and Eqns.(70-74) into Eqns. (75-79), we get the results as follows:

$$S(t) = a + (A - \mu a)t + (-\beta ac - A\mu + \mu^{2}a)\frac{t^{2}}{2!} + (\beta\mu ac - \beta Ac - \beta a(\alpha b - (\gamma + q + \mu)) + \mu(\beta ac + A\mu))\frac{t^{3}}{3!} + (\beta c(\beta ac + A\mu) + \beta A(\alpha b - (\gamma + q + \mu)) + \beta\mu Ac)\frac{t^{4}}{4!} + \beta^{2}Ac^{2}\frac{t^{5}}{5!}$$
(80)

$$E(t) = b - (\alpha + \mu)bt + (\beta ac + (\alpha + \mu)^{2}b)\frac{t^{2}}{2!} + (\beta Ac - (\alpha + \mu)\beta ac - \beta \mu ac + \beta a(\alpha b - (\gamma + q + \mu)))\frac{t^{3}}{3!} + (\beta A(\alpha b - (\gamma + q + \mu)) - \beta c(\beta ac + A\mu) - (\alpha + \mu)\beta ac)\frac{t^{4}}{4!} - \beta^{2}Ac^{2}\frac{t^{5}}{5!}$$

$$I(t) = c + (\alpha b - (\gamma + q + \mu)c)t + \left(-\alpha(\alpha + \mu)b + (\gamma + q + \mu)(\alpha b - (\gamma + q + \mu)c)\right)\frac{t^2}{2!} + \alpha\beta ac\frac{t^3}{3!} + \alpha\beta Ac\frac{t^4}{4!}$$
(82)

$$Q(t) = d + (qc - (\theta + \mu)d)t + (q(\alpha b - (\gamma + q + \mu)c) - (\theta + \mu)(qc - (\theta + \mu)d))\frac{t^2}{2!}$$
(83)

$$R(t) = e + (\gamma c + \theta d - \mu e)t + (\gamma (\alpha b - (\gamma + q + \mu)c) + \theta (qc - (\theta + \mu)d) - \mu (\gamma c + \theta d - \mu e))\frac{t^2}{2!}$$
(84)

Numerical Simulation and Discussions

We utilized the fourth-order Runge-Kutta technique to perform a numerical solution for the system of first-order nonlinear differential equations and MATLAB software was employed for this purpose. To gauge the accuracy of this numerical solution, a comparative analysis was conducted between these numerical results and the analytical solutions obtained through the Laplace Adomian Decomposition Method. Furthermore, we generated graphical representations of the analytical concentrations, namely S, E, I, Q, and R, along with their corresponding numerical results, covering a range of parameter values.

In the numerical testing phase, we adopt the following parameter values.
$$A = 1380, \beta = 0.0009, \alpha = 0.0037, \gamma = 0.0055, q = 0.00052, \theta = 1, \mu = 0.0028$$
 With the initial values $S(0) = a = 1, E(0) = b = 1, I(0) = c = 1, Q(0) = d = 1, R(0) = e = 1$

Figures 2-8 depict a comparison between numerical and analytical solutions, showcasing the concentration profile of SEIQR under varying sets of physical parameters. In Figure 2, we present the concentration of the Susceptible class, S(t), utilizing the specified parameters for both analytical solutions and numerical simulations, it occurs perfect fit. The Susceptible class is primarily influenced by the parameters A, β, μ with additional connections to other parameters through nonlinear equations. The influence of these parameters becomes evident as the concentration of the susceptible class demonstrates a notable increasing trajectory. Figures 3 and 4 illustrate the concentration profiles of the Exposed class, E(t), and the Infected class, I(t), respectively, employing the provided parameters for both analytical solutions and numerical simulations. The Exposed class and Infected class are primarily influenced by the parameters α, β, μ and α, γ, q, μ respectively, with additional connections to other parameters through nonlinear equations. The influence of these parameters becomes apparent as the concentration of the Exposed and Infected class exhibits a conspicuous upward trend. Figure 5 depicts the concentration profiles of the Isolated class, Q(t), utilizing the specified parameters for both analytical solutions and numerical simulations. The Isolated class is primarily influenced by the parameters q, θ, μ with additional connections to other parameters through nonlinear equations. The impact of these parameters becomes evident as the concentration of the Isolated class displays a noticeable decreasing trend. Figure 6 depicts the concentration profiles of the Recovered class, R(t), utilizing the specified parameters for both analytical solutions and numerical simulations. The Recovered class is primarily





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influenced by the parameters γ , θ , μ with additional connections to other parameters through nonlinear equations. The impact of these parameters becomes evident as the concentration of the Recovered class displays a noticeable increasing trend. Figure 7 portray the comprehensive concentration profile of SEIQR, encompassing both analytical solutions and numerical simulations. These figures showcase a satisfactory level of agreement within the time intervals t = [0,1].In Figure 8, we plot the behaviour of novel corona virus SEIQR model, by using the data set [10]: S(0) = 1380, I(0) = 121 and some values are assumed E(0) = 500, Q(0) = 70, R(0) = 40. In Figure 9, we plot the behaviour of novel corona virus SEIQR model, by assuming all data set as: S(0) = 380, E(0) = 150, I(0) = 121, Q(0) = 100, R(0) = 90.

CONCLUSION

This research is focused on obtaining an approximate analytical solution for the novel corona virus SEIQR model using the Laplace Adomian decomposition method (LADM). The LADM involves breaking down the equations into an infinite power series of Adomian polynomials and simplifying the computation of approximate solutions for the model's variables. We conducted a thorough comparison between the obtained analytical solutions and the numerical simulations. The findings demonstrated a notable level of consistency in the parameter values considered. The efficiency of the Laplace Adomian Decomposition method in handling nonlinear equations is underscored through graphical representation. Moreover, using different data set values, we analysed the behaviour of novel corona virus model. This comparative analysis will assist in the development of effective strategies to mitigate the spread of the virus and inform the implementation of appropriate public health interventions.

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Table 1: List of parameters used in this model.

Parameters	Description	
A	Recruitment rate	
β	Infection rate	
μ	Natural death rate and the disease death rate	
α	Rate at which exposed population moves to infected population	
θ	Rate at which isolated individuals recovered	
q	Turn Rate at which infected population moves to isolated population	
γ	Rate at which exposed population moves to isolated population	

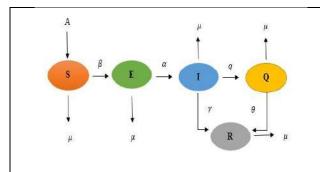


Figure 1: Flow chart of SEIQR model.

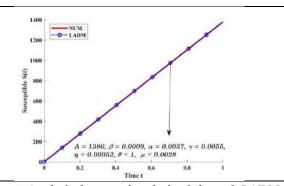


Figure 2: Analytical expression derived through LADM equation (80) is compared with numerical simulation for the concentration Susceptible S(t). The solid line corresponds to the numerical simulation, while *** represents equation (80).





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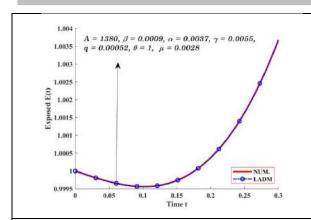


Figure 3: Analytical expression derived through LADM equation (81) is compared with numerical simulation for the concentration Exposed E(t). The solid line corresponds to the numerical simulation, while *** represents equation (81).

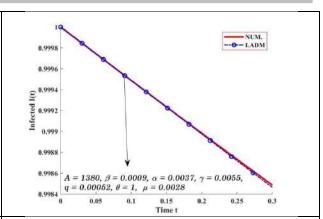


Figure 4: Analytical expression derived through LADM equation (82) is compared with numerical simulation for the concentration Infected I(t). The solid line corresponds to the numerical simulation, while *** represents equation (82).

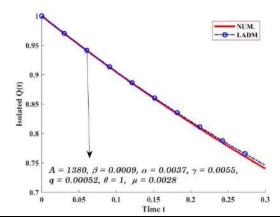


Figure 5: Analytical expression derived through LADM equation (83) is compared with numerical simulation for the concentration Isolated Q(t). The solid line corresponds to the numerical simulation, while *** represents equation (83).

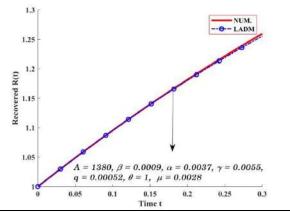


Figure 6: Analytical expression derived through LADM equation (84) is compared with numerical simulation for the concentration Recovered S(t). The solid line corresponds to the numerical simulation, while *** represents equation (84).





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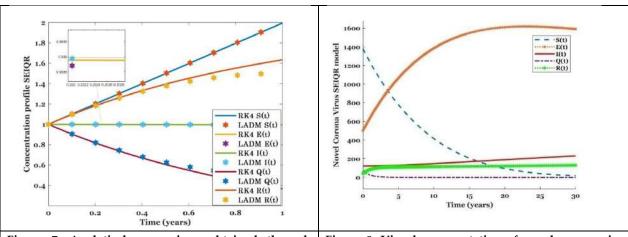
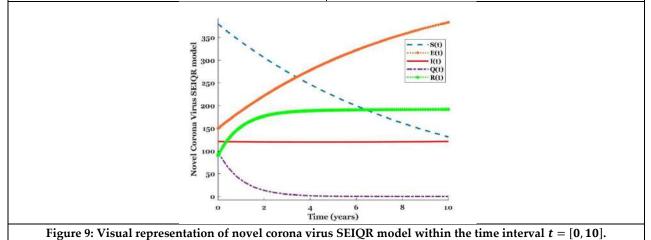


Figure 7: Analytical expressions obtained through LADM equations (80-84) are contrasted with numerical simulations for all SEIQR concentration profiles within the time interval t = [0, 1].

Figure 8: Visual representation of novel corona virus SEIQR model within the time interval t = [0, 30].







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RESEARCH ARTICLE

Evaluation of the Anti-Hyperlipidemic Potential of A Novel Anti-Diabetic Polyherbal Preparation in Streptozotocin-Induced Diabetic Rats and its Formulation in to a Capsule Dosage Form

Uppara Venkatesh^{1*} and B. V. Ramana²

¹Research Scholar, Department of Pharmacology, St. Johns College of Pharmaceutical Sciences, (Affiliated to Jawaharlal Nehru Technological University, Anantapur) Andhra Pradesh, India. ²Professor, Department of Pharmaceutics, Dr.K.V. Subba Reddy Institute of Pharmacy, (Affiliated to Jawaharlal Nehru Technological University, Anantapur) Andhra Pradesh, India.

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*Address for Correspondence Uppara Venkatesh

Research Scholar,

Department of Pharmacology,

St. Johns College of Pharmaceutical Sciences,

(Affiliated to Jawaharlal Nehru Technological University, Anantapur)

Andhra Pradesh, India.

Email: ukvenkatesh053@gmail.com



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ABSTRACT

Diabetes, also known as diabetes mellitus, is a long-term metabolic disease marked by high blood glucose levels. This condition is brought on by insufficient insulin production or poor insulin uptake by the body's cells. Diabetes can lead to hyperlipidemia, a condition marked by elevated blood triglyceride and cholesterol levels, as a result of the disturbance of lipid metabolism brought on by insulin resistance or insufficiency. In the current study, it was investigated whether a novel anti-diabetic polyherbal formulation made in-house could reduce hyperlipidemia in rats with diabetes induced by streptozotocin (STZ). The formulation contained dried leaves of Tinospora cordifolia, dried bark of Cinnamomum zeylanicum, dried seeds of Trigonella foenum, and dried seeds of Nigella sativa. Streptozotocin was injected intraperitoneally at a dose of 60 mg/Kg to induce diabetes in albino rats. Using STZ-induced diabetic rats, the effects of the Poly herbal preparation (PHP) were assessed for 28 days at doses of 200 and 400 mg/Kg body weight. Diabetic rats were treated with polyherbal preparation that was made inhouse, and the effects of the PHP on lipid metabolism were assessed by looking at various biochemical markers. The anti-hyper lipidemic efficacy was assessed using total cholesterol, triglycerides, very lowdensity lipoproteins, low density lipoproteins, and high-density lipoproteins. Since the ethanolic extract polyherbal formulation significantly reduced blood glucose levels in our earlier trials, it was selected for additional testing of its anti-hyperlipidemic potential. There were six rats in every group of thirteen. The blood glucose level was significantly (p<0.05) lowered upon oral administration of PHP extract (EPHP-F),





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with the effect becoming more pronounced at 400 mg/kg. There was dose-dependent action in the PHP. In comparison to normal, it significantly decreased the serum levels of triglycerides and cholesterol in diabetic rats. The outcomes verified that the PHP considerably decreased high blood cholesterol and triglycerides and had a dose-dependent blood glucose reducing impact. After that, a capsule formulation containing the optimal ratio of polyherbal preparation was formulated and evaluated for preformulation and quality control features. All polyherbal capsules were determined to have disintegration times that fell within acceptable pharmacopeial bounds. The resultant polyherbal capsule can be used as a stable, patient-friendly, and cost effective solid dose form.

Keywords: Lipid, triglycerides, cholesterol, Tinospora cordifolia, Cinnamomum zeylanicum, Trigonella foenum, Nigella sativa.

INTRODUCTION

The impact and affection that diabetes mellitus (DM) has caused throughout the world in recent decades has been worrisome. There were 463 million diabetics worldwide in 2019, and by 2045, that number is expected to rise to 700 million. Diabetes mellitus (DM) is a chronic degenerative illness linked to high blood sugar levels that damages systems, organs, and cells. Overabundance of blood glucose leads to the production of reactive oxygen species, which in turn causes oxidative stress to escalate. This stress can result in the formation of advanced glycation end products (AGEs), dyslipidemia, and various metabolic disorders that are associated with the illness. Diabetes can manifest in various forms, one of which is type 1 diabetes (DM1), a condition brought on by an insulin-dependent lifestyle. Additionally, there is type 2 diabetes (DM2), which is the most common kind and is linked to elevated glucose concentrations as well as insulin sensitivity or resistance. Microvascular issues (retinopathy, nephropathy, and neuropathy) and macrovascular issues (ischemic heart disease, peripheral vascular disease, and cerebrovascular disease) are closely related in roughly one-third to one-half of DM patients. Unchecked hyperglycemia can lead to oxidative DNA damage by tissue protein glycation, enhanced lipid peroxidation, superoxide production, and lipoprotein glycation. Glycation and glucose auto-oxidation (OH-) produce reactive species like hydroxyl radicals and hydrogen peroxide (H2O2). It consequently affects human life globally in one or more ways. [5-6] DM is complicated and challenging to manage.

Several medical interventions are used to try to control blood glucose levels. Research is always ongoing to find novel options that can slow the progression of diabetes diseases.[7] Even though a number of conventional oral hypoglycemic drug types have been introduced to the market, the prevalence of diabetes and health care costs are rising alarmingly globally. The global economy is predicted to suffer losses due to diabetes in 2019, 2030, and 2045, of USD 760 billion, USD 825 billion, and USD 845 billion, respectively. [8] These problems therefore call for the creation of novel antidiabetic drugs that are structurally different from those that are currently on the market, safe, easily obtainable, more effective, and have a novel mode of action. These drugs should also be derived from natural sources, mainly plants.[9] Due to their favorable biological actions, ethanobotanical extracts of medicinal plants have been examined for their richness in phytochemical substances. These extracts have historically been utilized as therapeutic systems.[10] Because they can offer a more all-encompassing and synergistic approach to the condition, poly herbals—formulations made up of a variety of different herbs and plant-based ingredients—have become more popular in the management of diabetes. Diabetes is a complex metabolic disease that affects several physiological processes, such as glucose metabolism, insulin control, and oxidative stress. Herbs that target various elements of diabetes are frequently included in polyherbal medicines with the goal of enhancing insulin sensitivity, improving blood sugar regulation, and providing antioxidant protection. Poly herbals, which include these herbs, can provide a comprehensive approach to managing the many aspects of diabetes, possibly yielding greater advantages than the use of individual herbs. [11-12]





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Poly herbals have a significant role in managing diabetes since they can help people achieve better glucose control while reducing the negative effects of some drugs. [13] But it's important to remember that there are differences in the safety and effectiveness of polyherbal supplements, therefore use should be supervised by a doctor. To determine the efficacy of certain formulations, the best dosages, and any possible interactions with traditional diabetic therapies, clinical research is required. Healthcare providers should provide assistance when including polyherbals into a diabetic management strategy, as they should be viewed as an adjunct to prescribed drugs and lifestyle adjustments. This method emphasizes the need of individualized, patient-centered care in achieving the best possible outcomes for people with diabetes while acknowledging the complex nature of the disease. [14-15] In previous studies, we prepared n-hexane, petroleum ether, and ethanol extracts from a poly herbal preparation made up of different amounts of dried Trigonella foenum (TF) leaves, dried Nigella sativa (NS) seeds, dried bark of Cinnamomum zeylanicum (CZ), and dried leaves of Tinospora cordifolia (TC). Using DPPH (2,2-diphenul-1-picrylhydrazyl) and hydrogen peroxide scavenging assays, we evaluated the in vitro antioxidant characteristics of these extracts; the ethanolic extract showed significant antioxidant activity. Additionally, we looked at the anti-diabetic properties of ethanol, petroleum ether, and n-hexane extracts at 200 and 400 mg/KG body weight. Of these, the polyherbal preparation's ethanol extract (EPHP 1), which contained TC, CZ, TF, and NS in a 1:1:1:1 ratio, demonstrated a noteworthy decrease in blood glucose levels that was on par with glibenclamide, a common medication. Building on these discoveries, we subsequently conducted studies to determine the anti-hyperlipidemic potential of EPHP 1 in diabetic rats caused by streptozotocin and to formulate PHP into an appropriate dose form. [16]

MATERIALS AND METHODS

Chemicals

We bought streptozotocin (STZ) from Merck in Mumbai, India. We bought petroleum ether, n-hexane, and AR grade ethanol from SD Fine Chem in Mumbai, India. Glibenclamide (GLB) was graciously provided by Dr. Reddy laboratories in Hyderabad, India. Analytical grade chemicals were all that were employed in this investigation.

Plant material

We bought dried *Nigella sativa* seeds, dried *Trigonella foenum* seeds, and dried *Cinnamonum zeylanicum* bark from the local market in Kurnool, Andhra Pradesh, India. From its natural environment in and around the Nallamala Forest area near Srisailam, Kurnool District, Andhra Pradesh, India, the dried leaves of *Tinospora cordifolia* were collected. The plants were verified by Dr. K. Madhava Chetty, Associate Professor in the Department of Botany at Sri Venkateswara University in Tirupathi, Andhra Pradesh, India. Parts of plants were shade dried.

Preparation of extract

Using n-hexane, petroleum ether, and ethanol as solvents, the dried leaves of TC, dried bark of CZ, dried seeds of TF, and dried seeds of NS were continuously extracted using a soxhlet apparatus. First, each of the 2.5 kg of plant components was air-dried and then independently ground into a coarse powder. Following meticulous weighing, 1 kg of each crude drug powder was added to a soxhlet system and extracted for 48 hours at a temperature between 70 to 80 °C using the previously indicated solvents. Dark brown to black extracts were produced by continuing the extraction process until the solvent was clear. The extracts were cooled and filtered to remove any remaining material. Using a rotary evaporator, the extracts were condensed under low pressure to create a powder, which was subsequently dried. After calculating the extract percentage, the extracts were refrigerated and kept in amber glass containers for later processing. To determine which phytoconstituents were included in the extracts, preliminary phytochemical studies were carried out. In order to conduct the investigation, the dry powder was diluted with the necessary amount of 0.5% carboxymethyl cellulose (CMC).

Animals

Both male and female albino Wistar rats weighing 160–200 grams were purchased from SV Animal House and Enterprises in Bangalore, India. These rats were used for assessments of their anti-diabetic activity as well as acute





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toxicity investigations. The animals went through a 10-day stabilization period following procurement. They were kept in cages made of polypropylene with a 12-hour light-dark cycle, 60±5% relative humidity, and room temperature. The rats were fed a regular pellet diet and had unlimited access to water during the trial. The animals were handled carefully to reduce any potential discomfort that may otherwise lead to an increase in adrenal output.

Development of polyherbal preparation

By combining the dried extracts of the plant components in different ratios, the polyherbal preparation (PHP) of the dried leaves of TC, dried bark of CZ, dried seeds of TF, and dried seeds of NS was created. Table 1 displays the PHPs' makeup.

Dosage selection and preparation

Two doses were selected based on the acute toxicity experiments in order to evaluate the antidiabetic activity. There were two doses: 200 mg/kg and 400 mg/kg body weight; the first dose was roughly one-tenth of the maximum dose of 2000 mg/kg, while the second dose was twice that much. To prepare the 200 mg/kg and 400 mg/kg doses of the polyherbal preparation (PHP), the required number of extracts were dissolved in a 0.5% sodium CMC solution.

Animal grouping

There were 13 batches of animals total, with 6 animals in each group (n = 6). Regular control diabetic control, Glibenclamide treated groups and PHP treated groups at 200 mg/Kg(EPHP-A to EPHP-E) and groups treated with polyherbal preparation at doses of 400 mg/kg (EPHP-F to EPHP-J). For 28 days, the therapy was administered nonstop.

Induction of Diabetes

The animals were given a single intraperitoneal injection of freshly prepared streptozotocin (STZ) at a dose of 60 mg/kg to induce diabetes. Prior to injection, the STZ was dissolved in regular saline. After the animals had the STZ injection, they were watched for a whole day. Following this initial phase, the animals received a 10% glucose solution for a full day. The blood glucose level (BGL) was measured on the third day in order to verify the diagnosis of diabetes. This assessment acted as a gauge to see if the animals' diabetes was successfully brought on by the STZ injection.

Determination of antidiabetic activity

Oral gastric gavages were used to provide the test samples to the animals once a day, before they were allowed to eat. At the start of the trial, the animals' blood glucose levels were determined using a glucometer. The 7th, 14th, 21st, and 28th days after the experiment's beginning saw a recurrence of these measurements. This made it possible to assess the test samples' possible effects on blood glucose levels over the course of 28 days.

Estimation of biochemical variables

The animals were sacrificed by cervical dislocation on day 28, and the biochemical variables were assessed. Using an auto-analyzer, the glucose oxidase method was used to analyze total cholesterol (TC), triglycerides (TG), low density lipoproteins (LDL), high density lipoproteins (HDL), and very low-density lipoproteins (VLDL).

Formulation of a poly herbal capsule

The factual extracts and excipients were all run through British Standard Sieves (BSS) #120. Using an electronic balance, a polyherbal preparation that was tailored for both performance and safety was precisely tested. The mixture was run over BSS # 40 once more, mixed twice, and then put into size one capsules to create granules. An evaluation was conducted on the preformulation parameters, including the hausner's ratio, bulk and tapped density, compressibility index, and angle of repose.





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An analysis of the capsules

The Electrolab dissolving apparatus was utilized to conduct the disintegration test. Six baskets were filled with one capsule each, and the device was maintained at 37±0.5°C of the immersion liquid. The amount of time it took for the capsule to break down totally was noted. The capsules dissolve when, after going through a 10# mesh screen, no particles are visible above the gauge.

Analytical statistics

The mean \pm Standard error mean (SEM) was used to express the study's data. A one-way analysis of variance (ANOVA) was used to analyze antidiabetic behavior, anti-hyperglycemic effects, and other pertinent data. Dunnet's Multiple Comparison Test was then employed to confirm the results. A statistical significance threshold of P < 0.05 was applied to ascertain the results. P values less than this cutoff were regarded as statistically significant, signifying a meaningful difference or effect in the results.

RESULTS

PHP's impact on the blood glucose levels of diabetic rats

When compared to the negative control group, the diabetic control rats' blood glucose levels increased gradually and consistently over the course of our experiment, and this difference was statistically significant (P<0.05). Nevertheless, after the rats received GLB (5 mg/kg) and PHPs, there was a significant drop in blood glucose levels beginning on the seventh day after the rats were exposed to these medications (P<0.05). This effect persisted for the duration of the study, lasting until day 28, when the PHPs' drop in blood glucose levels was even more significant (P<0.05), outperforming the effectiveness of GLB, the standard drug. Table 2 lists the anti-diabetic effects of ethanol extracts, particularly EPHP, given at doses of 200 and 400 mg/kg of body weight. These findings highlight PHPs' enormous potential for controlling blood glucose levels in diabetic settings. PHPs have shown promise as a valuable therapeutic option for diabetes, as seen by the prolonged reduction in blood glucose levels and their superior effectiveness on day 28 when compared to the conventional medicine. The entire scope of their use in the treatment of diabetic patients may become clear through additional investigation and clinical trials.

Impact of EPHP on diabetic rats' lipid profiles

When compared to normal rats, there was a significant (P<0.001) increase in serum levels of TC, TG, VLDL, and LDL and a significant (p<0.001) drop in HDL cholesterol following the creation of diabetes and subsequent treatment with the two doses (200 and 400 mg/KG) of PHPs. When compared to the diabetic control, the animal groups treated with PHP at a dose of 200 mg/KG body weight (EPHP -A to EPHP -E) did not exhibit a significant decrease in the levels of TC, TG, LDL, or VLDL, nor did they exhibit a significant increase in the levels of HDL in the serum. Thus, the blood levels of TC, TG, LDL, and VLDL were significantly lower in the animal groups that received a therapy of PHP at 400 mg/KG body weight (EPHP -F to EPHP -J). There was a noticeable increase in the serum HDL level as compared to the diabetes control group. When EPHP (200 and 400 mg/KG) was given to diabetic rats for 28 days, the serum levels of TC, TG, LDL, and VLDL were significantly (p<0.001) reduced, but the levels of HDL were increased in comparison to the diabetic control group. Table 3 displays the findings of the impact of EPHP on the lipid profiles of diabetic rats.

Analysis of polyherbal capsules

Preformulation testing was performed on the optimized PHP i.e., EPHP-F powder mixture. The mix was evaluated for bulk density, tapped density, angle of repose, compressibility index, and Hausner's ratio in order to analyze flow behavior. When the mixture was enclosed in a capsule, its flow properties were satisfactory. This is confirmed by the data in table 4. It was discovered that the capsules disintegrated on average in 13.6±1.07 minutes. The polyherbal capsules are depicted in Figure 1.





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DISCUSSION

Medicinal plant parts are widely used by humans to treat a wide range of illnesses, including diabetes, even though there isn't enough scientific data to support their safety and effectiveness. Therefore, it is essential to thoroughly evaluate the safety and effectiveness profile of medicinal plants used in conventional medicine. [17] Due to the limitations of traditional anti-diabetic medications, there is an urgent need for new, more effective, affordable, and nontoxic medications. There is a great deal of demand for natural product-based alternative medicines since they are regarded as safe, readily available, and do not require the lengthy manufacturing of pharmaceuticals. [18] It has long been known that polyherbal formulations have great promise for the treatment of a broad range of illnesses and ailments. The inclusion of a wide variety of phytoconstituents that work in concert is thought to be responsible for these formulations' medicinal efficacy. Interestingly, these formulations show good performance even at lower dosages and continue to be safe at higher concentrations.[19] Our study focused on evaluating the hepatoprotective and nephroprotective qualities of an in-house made ethanolic extract of a novel polyherbal mixture made up of four different plant extracts because of the therapeutic potential of polyherbal formulations. [20] Dried leaves of *Tinospora cordifolia*, dried bark of *Cinnamomum zeylanicum*, dried seeds of *Nigella sativa*, and dried seeds of *Trigonella foenum* were among these extracts. Rats with diabetes caused by streptozotocin (STZ) were used in this study. [16]

We carried out an investigation into the acute oral toxicity of our earlier trials. Administration of the curd test extract at a single limit test dose of 2000 mg/KG did not result in mortality or delayed toxicity throughout the 14-day followup period. The results showed that there is a significant margin of safety and that the test extracts LD50 value is anticipated to be greater than 2000 mg/KG. Overall, the findings demonstrated the palatability and nontoxic nature of EPHP, suggesting that the plant is suitable for usage in conventional settings. [16] It is believed that several phytochemical components that have been isolated from different plants have strong hypoglycemic, antihyperglycemic, and glucose-lowering properties. Alkaloids, triterpenoids, phenolics, and flavonoids are a few of the bioactive components. The action may be accompanied by increased pancreatic β-cell release of insulin, decreased small intestine absorption of glucose, increased body consumption of glucose, and/or the initiation of hepatic glycogenesis. Furthermore, it has been amply demonstrated that these potent phytochemical substances may repair damaged beta cells in diabetic rats while protecting them from oxidative stress. [21] The most widely used and accepted model of experimental diabetes in rats is streptozotocin-induced diabetes. This diabetogenic drug is generally preferred over alloxan due to its wider species effectiveness and reproducibility. According to published research, rats can experience chronic hyperglycemia from the inducer after a single dose of 60 mg/kg for at least a month. [22] Similar to diabetic management, at this dosage the chemical results in chronic hyperglycemia with no appreciable change in blood glucose levels for two weeks.

To assess the impact of the extracts on BGLs, normoglycemic and STZ-induced diabetic rat models were established. The vehicle-treated groups did not show a statistically significant decrease in fasting blood glucose levels in the model of normoglycemic rats. Over the course of 28 days, blood glucose levels in diabetic rats significantly increased. When compared to the standard medication glibenclamide, the diabetic rats treated with 200 mg/KG body weight (EPHP-A to EPHP-E) demonstrated a reduction in BGL, but the efficacy was noticeably lower. On the other hand, the BGL was significantly lower in the diabetic rats who received EPHP at a dose of 400 mg/KG body weight (EPHP-F to EPHP-J). Especially EPHP-F, which combines TC, CZ, TF, and NS at a ratio of 1: 1: 1: 1, when administered at a 400 mg/KG body weight, exhibits a significant decrease in BGL in comparison to the standard drug group. The maximum drop in blood glucose in STZ-induced diabetic rats was seen on day 28 at 400 mg/KG of extract, which was equivalent to a normal dose. It was noted that a dose-dependent relationship existed between the test extract activity. This may indicate that the PHP's bioactive ingredients, which are essential for its anti-diabetic effects, are more concentrated at greater dosages. Accordingly, the present study hypothesized that in a time- and dose-dependent way, the ethanol extract of PHP, in particular EPHP-F, has exceptional anti-diabetic action on diabetic rats caused by STZ. The results imply that the test extract might have a strong hypoglycemic impact. This conclusion is supported by logical arguments. It has been suggested that by lowering oxidative stress, using antioxidants can help





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reduce diabetic complications. The main factors contributing to the development and course of diabetes are inflammation and oxidative stress. Because STZ selectively destroys pancreatic beta cells, it results in diabetes mellitus. Research indicates that when administered to animals, it causes \(\beta cells \) to die after three days. Results showed that PHP containing dried Nigella sativa seeds, dried Bark of Cinnamomum zeylanicum, dried Leaves of Tinospora cordifolia, and dried Seeds of Trigonella foenum exhibited strong antioxidant activity. This suggested that the β-cells' ability to guard against oxidative free radicals may be the cause of the anti-hyperglycemic activity in diabetic rats induced by STZ. The release of insulin from pancreatic beta cells may have been the PHP extract's mode of action. The anti-diabetic benefits of the ethanolic extract of PHP may be explained by the presence of many bioactive components that have been linked to hypoglycemic activity. In individuals with diabetes, hypertriglyceridemia and hypercholesterolemia are the most prevalent lipid abnormalities. [23] High serum TC, TG, LDL, VLDL, and low HDL values are its defining characteristics. Low insulin causes hormone-sensitive lipase to be activated, which increases the liver's production of VLDL and lipolysis. [24] Additionally, lipoprotein lipase activity is decreased in insulin insufficiency, which may result in a decrease in the clearance of VLDL and chylomicrons. [25] Over the course of 28 days, the administration of EPHP dramatically decreased levels of TC, TG, LDL, and VLDL while dose-dependently raising HDL levels. This result was consistent with prior findings that demonstrated the ethanol extract's strong antidyslipidemia and anti-hyperglycemic effects. The observed antihyperlipidemic impact may be explained by decreased chelosterogenesis and fatty acid synthesis caused by inhibition of pancreatic lipase and pancreatic cholesterol esterase, respectively. These results highlight EPHP-F's promising potential in the context of its antidiabetic and anti-hyperlipidemic effects. They also demonstrate the drug's ability to prevent the progression of hyperlipidemia and mitigate lipid-related damage, which could have a major positive impact on the management of lipid-related complications. To further investigate its clinical applicability in this particular situation, more clinical research is necessary.

CONCLUSION

The EPHP-F had a positive impact on blood glucose levels in this investigation. Additionally, it raised HDL levels and corrected serum triglycerides and cholesterol. The study's findings demonstrated that EPHP-F has potent anti-diabetic and anti-hyperlipidemic properties, matching or surpassing those of glibenclamide, a common medication. The study's findings might be applied to the creation of novel anti-diabetic medications for the control and therapy of type 2 diabetes. To determine the molecular mechanism underlying its anti-diabetic and anti-hyperlipidemic effects, more investigation is needed to pinpoint the molecular targets and isolate the bioactive phytoconstituents.

Ethical endorsement

The study technique SJCP/PCOL/AD2022-10/011 was approved by the institutional animal ethics committee. Under the registration number 1519/PO/Re/S/11/CPCSEA, the committee for the regulation and supervision of animal experimentation made recommendations, and the ethics committee followed those suggestions in conducting the studies.

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Table 1 Composition of polyherbal preparations

Extracts	PHP Code	PHP formulation	Ratio
	EPHP 1	TC: CZ: TF: NS	1: 1: 1: 1
Eth an al autus at	EPHP 2	TC: CZ: TF: NS	2: 2: 2: 1
Ethanol extract	EPHP 3	TC: CZ: TF: NS	2: 2: 1: 2
	EPHP 4	TC: CZ: TF: NS	2: 1: 2: 2
	EPHP 5	TC: CZ: TF: NS	1: 2: 2: 2

Table 2 Anti diabetic activity of Ethanol extracts of PHP on STZ induced diabetes in rats

Carona	Tuestmeent	Blood glucose levels (mg/dL) on					
Group	Treatment	0 day	7 th day	14 th day	21st day	28th day	
Normal control	Vehicle (0.5 % Na CMC)	76.68 ± 2.03	78.36 ± 2.24	82.45 ± 2.02	80.11 ± 1.86	84.77 ± 2.33	
Diabetic	STZ 60 mg/Kg i.p	78.29 ±	285.16 ±	292.11 ±	304.98 ±	313.73 ±	
control	312 60 Hig/Kg 1.p	3.23 a	3.82 a	4.12 a	4.85 a	4.36 a	
Standard	STZ 60 mg/Kg i.p+ GLB 5	73.28 ±	267.79 ±	202.15 ±	166.62 ±	121.36 ±	
Standard	mg/Kg	3.65 в	3.94 в	3.54 b	4.17 c	4.21 ^c	
EPHP -A	STZ 60 mg/Kg i.p+ EPHP 1	70.71 ±	277.84 ±	213.99 ±	179.67 ±	167.28 ±	
EFFIF -A	200 mg/Kg	5.32 °	4.59 b	6.13 c	3.80 c	4.01 c	
EPHP -B	STZ 60 mg/Kg i.p+ EPHP 2	74.38 ±	282.29 ±	195.74 ±	179.61 ±	166.12 ±	
ЕГПГ-В	200 mg/Kg	7.91 b	5.22 b	3.82 c	4.76 b	4.28 b	
EDLID C	STZ 60 mg/Kg i.p+ EPHP 3	79.65 ±	275.41 ±	215.03 ±	176.21 ±	163.95 ±	
EPHP -C	200 mg/Kg	7.72 °	5.86 c	5.04 b	5.14 b	5.62 °	
EDITO D	STZ 60 mg/Kg i.p+ EPHP 4	68.76 ±	286.55 ±	212.89 ±	174.42 ±	162.14 ±	
EPHP -D	200 mg/Kg	3.19 ^b	3.45 °	3.58 b	4.02 b	3.97 °	
EDIID E	STZ 60 mg/Kg i.p+ EPHP 5	78.78 ±	283.62 ±	223.40 ±	200.94 ±	165.15 ±	
EPHP -E	200 mg/Kg	4.31 °	3.66 c	2.87 b	4.18 b	3.99 ь	
EDITO E	STZ 60 mg/Kg i.p+ EPHP 1	66.94 ±	268.78 ±	205.63 ±	157.12 ±	126.42 ±	
EPHP -F	400 mg/Kg	6.11 °	4.95 °	4.62 ^c	4.25 b	4.05 b	
EDLID C	STZ 60 mg/Kg i.p+ EPHP 2	67.79 ±	283.12 ±	201.65 ±	164.89 ±	133.57 ±	
EPHP -G	400 mg/Kg	5.17 °	4.32 b	4.66 c	3.87 ^c	4.08 b	
EDIID II	STZ 60 mg/Kg i.p+ EPHP 3	65.16 ±	280.02 ±	215.28 ±	170.11 ±	136.49 ±	
EPHP -H	400 mg/Kg	7.49 ^b	6.54 b	5.89 °	5.82 ^c	5.16 °	
EDIID I	STZ 60 mg/Kg i.p+ EPHP 4	61.78 ±	281.02 ±	201.46 ±	188.93 ±	143.21 ±	
EPHP -I	400 mg/Kg	6.15 c	4.93 b	4.32 ^c	4.14 ^c	3.56 c	
EDIID I	STZ 60 mg/Kg i.p+ EPHP 5	66.17 ±	282.41 ±	206.94 ±	185.03 ±	136.82 ±	
EPHP -J	400 mg/Kg	6.28 b	4.78 c	5.02 b	5.41 b	4.68 b	





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Values are expressed as mean \pm SEM., Data analyzed by one way ANOVA followed by Dunnet's Multiple Comparison Test. a-p<0.05 as compared with control group, b-p<0.01 and c-p<0.05 as compared with STZ group.

Table 3 Effect of EPHP on lipid profile

Caracan	Tuestuesut		Lipid p	rofile (mg/dL) on	
Group	Treatment	TC	Triglycerides	VLDL	LDL	HDL
Normal control	Vehicle (0.5 % Na CMC)	81.55±2.67	54.73±1.45	16.34±1.45	46.47±0.79	39.78±1.92
Diabetic control	STZ 60 mg/Kg i.p	211.74±1.69	138.89±2.80	65.78±2.78	111.45±1.35	7.55±1.78
Standard	STZ 60 mg/Kg i.p+ GLB 5 mg/Kg	101.64±2.18°	77.54±1.56 °	23.79±2.78 c	52.23±1.78 °	30.74±1.34
EPHP -A	STZ 60 mg/Kg i.p+ EPHP 1 200 mg/Kg	135.75±2.67	92.78±2.78 °	42.39±3.73	74.94±1.89°	19.45±2.84
ЕРНР -В	STZ 60 mg/Kg i.p+ EPHP 2 200 mg/Kg	130.64±1.34	86.89±2.89 °	46.55±2.10	71.56±1.78°	15.48±1.69
ЕРНР -С	STZ 60 mg/Kg i.p+ EPHP 3 200 mg/Kg	131.54±2.34	91.42±2.75 °	41.89±2.38	70.34±1.89°	11.47±.02 °
EPHP -D	STZ 60 mg/Kg i.p+ EPHP 4 200 mg/Kg	126.64±2.34	94.56±2.87 °	43.64±1.54	71.54±2.67 °	14.56±2.89
ЕРНР -Е	STZ 60 mg/Kg i.p+ EPHP 5 200 mg/Kg	130.54±1.73	89.22±1.31 °	46.85±1.56	74.50±1.90 °	14.70±2.09
EPHP -F	STZ 60 mg/Kg i.p+ EPHP 1 400 mg/Kg	105.64±1.78	80.77±4.11 °	29.58±7.87	56.33±5.87°	32.46±8.45
EPHP -G	STZ 60 mg/Kg i.p+ EPHP 2 400 mg/Kg	116.69±1.78	89.68±2.92 °	31.64±2.84	57.74±1.23 °	29.67±2.84
EPHP -H	STZ 60 mg/Kg i.p+ EPHP 3 400 mg/Kg	120.64±1.99	94.67±2.90 °	37.54±0.34	59.79±0.89°	30.87±1.43
EPHP -I	STZ 60 mg/Kg i.p+ EPHP 4 400 mg/Kg	119.53±1.67	86.57±2.78 °	37.45±2.14	62.34±1.33 °	28.34±1.23
EPHP -J	STZ 60 mg/Kg i.p+ EPHP 5 400 mg/Kg	120.57±1.58	83.49±2.52 °	32.80±2.48	58.84±2.34 °	29.33±2.40

Values are expressed as mean ± SEM., Data analyzed by one way ANOVA followed by Dunnet's Multiple Comparison Test. a-p<0.05 as compared with control group, b-p<0.01 and c-p<0.05 as compared with STZ group.

Table 4 Preformulation evaluation of EPHP-F blend

Trail	Flow Properties				
1 raii	Angle of repose Bulk density		Tapped density	Compressibility index	Hausner's ratio
1	25.78	0.73	0.71	5.86	1.01
2	27.14	0.65	0.63	6.54	1.04
3	26.02	0.70	0.74	5.24	1.01
4	25.68	0.64	0.66	6.82	1.03
5	27.46	0.71	0.72	5.61	1.06
6	26.91	0.69	0.68	7.73	1.04
Mean	26.498±0.697	0.687±0.031	0.690±0.037	6.300±0.833	1.032±0.0177





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Figure 1 Photograph of capsules





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REVIEW ARTICLE

An Innovative Review about Audio and Video Collaboration for Event **Detection using Deep Learning Approaches**

M.Rajamanogaran^{1*} and G. Karthikeyan²

¹Ph.D Research Scholar, PG and Research Department of Computer Science, Periyar Government Arts College, Cuddalore, (Affiliated to Thiruvalluvar University, Vellore) Tamil Nadu, India.

²Assistant Professor, PG and Research Department of Computer Science, Periyar Government Arts College, Cuddalore, (Affiliated to Thiruvalluvar University, Vellore) Tamil Nadu, India.

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*Address for Correspondence

M.Rajamanogaran

Ph.D Research Scholar, PG and Research

Department of Computer Science,

Periyar Government Arts College, Cuddalore, (Affiliated to Thiruvalluvar University, Vellore)

Tamil Nadu, India.

Email: manogaran248@gmail



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ABSTRACT

Another milestone in the outbreak of artificial intelligence is the event detection done automatically by the machine in the visualization of human eye. In current years event detection is catching popularity and gaining awareness due to rapid growth of online videos in various fields like sports, entertainment, education, flash news etc. Event detection similar to object recognition is the method used to identify the location of the entities thereby trying to describe the scenery in the video. This process is enabled by two measures like audio and video to strengthen the depiction. The correct mixing of audio along with the video scenes elaborates the details of the displays. The precise method used is the deep learning approach where the event detector learns the occurred actions in the particular scene by studying the features and classifying according to the predicted classes. This paper is the general survey about the methods used for acquiring the data sets and the medium for processing by using different feature extraction and classification algorithms.

Keywords: event detection, audio, video, deep learning, features, classifying, survey, feature extraction, classification.





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INTRODUCTION

The latest development in image processing and computer insight leads to the improved version of object detection as event recognition to deal with identification of variety of objects that belong to certain section. In digital images and videos the actions performed are narrated along with the picture automatically. This is possible due to the rapid enhancement in the digital world along with the network facility that brought the attentiveness among social media. [1]. Artificial intelligence plays its role in copying the human activities to be performed by machines by training with sample data and testing for accuracy. One of these applications is the event detection method to predict the result by analysing the actions in the particular video [8] scene along with the captured audio for detecting the extract. The induced use of social media in various fields encourages the development of event detection as a demanding research zone to exploit the challenges for further use. It is defined as the process of verifying the streams of actions to identify the equivalent designs to produce the output as the required event. The types are specified by the backgrounds and patterns [9-13] of the event. The particular features of the objects like shape, texture is used for recognitions of the events and the user should be careful with problems like over fitting, occlusion, reacquisition and blurring of images while performing event detection. Event detection contains video identification where the videos are converted into frames of images and the frames are classified into different groups based on the actions performed in the video. In videos the objects are identified on the basis of video tracking tasks and object identification [2] by joining the advantages of both the methods into a mutual combined route.

The video contains the actions with properties like region, period, abode which takes place either normally or artificially. Identifying such events automatically is the challenging task for researchers. Displayed and online videos are embedded with audio produced in the sight which is also used for tracking the scene in the video. Sound is used for defining the activities that are produced in the videos by appreciating the circumstances of the event. Sound also identifies the actions that matches with predefined events by using the theme of the actions performed. Due to the presence of memory chips for sound and digital processing unit for audio [3], tracing of sound is simplified but the background overlapping of noise may reduce the quality of audio that should be avoided in the event detection. This can be achieved by acquiring the signals continuously and changing into symbols for explanation of the actions performed. Various applications include multimedia events, health care, surveillance and in other research areas. Event detection faced many issues when implemented with machine learning models and algorithms for huge data set with increased computational power. So deep learning models gained reputation due to their advanced techniques. Accordingly [4] novel neural network models are developed for accurate results to foretell the events in the video scene. Deep [14] convolutional network and recurrent neural network with unsupervised features are proposed to classify multi labelled data in event detection. Deep learning performs well with large data sets [15-18] and training of data needs abundant sample for accurate results.

APPLICATIONS IN REAL WORLD

This section provides some applications using video and audio [19] tracking for event detections in real life incidents. There are separate applications for both audio and video. Some are described in brief. Video applications in event tracking There are some presentations that make use of videos [25-30] for event tracking recently. Some of them are described in short-term.

Video Surveillance

Nowadays cameras are allowed to be used in offices and factories to monitor the workers and visitors to avoid accidents and any malpractice that may happen. Video tracking is used in [42] such systems to detect any objects that are present in the scenes. Many shops and crowded areas like hospitals, malls, theatres etc. use CCTV cameras for surveying purpose [68].





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Facial identification

Recognition of individual face is one of the types of biometrics[36][37] used as password in locking of smartphones and houses for safety purposes. Video tacking for object detection is also used for entertainment commitments using filters in many applications like Instagram, YouTube etc.

Self-Driving cars

Development in technology has introduced self- driving cars making use of video tracking [64-67] for event detections. The systems using video [31-36] tracking can detect active and passive items along the path to locate moving passengers and objects so that they can maintain the correct lane. Some examples are traffic lights, foottravelers using automatic event detection with the help of videos so that accidents could be avoided.

Audio applications in event detection

Recently many researches are taking place in the field of audio [20-24] applications in event tracing. The solicitations that are familiar are discussed below.

Surveillance

In monitoring incidents [63] sound effects also help in detecting features along with vision. In activities like accident, assassination, shooting etc. sound performs well for surveillance of the events more accurately. Many multimedia applications use audio structures for classification of ordinary and unusual events using several audio features. [5]. Audio deals with noisy situations to predict different sounds to match with the incidents happened in that environment.

Meeting

During crowded places like meeting, [38] social get together, parties, functions audio plays important role. Appreciation, encouragement can be shown in sound media like clapping, whistle sounds for positive and any rude incidents can be clearly shown in audio effects like unusual high decibel noise. Instead of body language audio can also describe the scenery [39] that takes place in the particular video that focuses on the meeting. Audio is also used for comparing the performance of various models in event detection. While dividing the video scenes into different segments audio information is used for accuracy.

Sports Events

Sports play a vital role in everyday activities. Observing highlights from the sports events needs audio effects also along with video for event identification. While recording the game audio tracing is also needed for effective visualization. So, research has invented many models for tracking the sequences of audio in the sports. Games like tennis, football require sound and models like Markov, Gaussian Mixture Model are used for segmentation of audio into various sequences for better understand ability.

Entertainment

Entertainment media like movies, short films, videos[40], online games make use of audio tracking to represent action scenes like robbery, explosion, flood and violent scenes like murder, gangster activities. Familiar networks like Bayesian model are used for obtaining information about audio segments to detect peculiar sound in the video scenes. Individual audio words are described by the segments for various characteristics.

Consumer Video

Various methods are proposed for classification using audio properties in consumer video. Each video is generated as frames and they are classified into multi actions based on the sound. The video features are extracted and mixed with audio types to detect particular event in the clippings using frequency analysis along with wavelet coefficients. There are various applications that make use of audio and video [41] for event tracking in different media. Some of them are focussed in this paper. Research work is carried in this field for further proceedings.





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DATA SET DESCRIPTION

There are numerous collections of data set for both video and audio as free source available for access. The collection contains both the copies of sound and picture along with the contributors and contains the description of the saved files like scope and format. Some data sets are discussed for both audio and video files.

Audio Data set

One of the popular data sets is Free Spoken Digit containing 3000 recordings of size 10Mb discussing about spoken English digits. Another standard one is, The spoken Wikipedia Corpora with 5397 collections of 23Gb size. It enumerates aligned spoken articles. Speech Commands Data set contains a maximum of 65000 tracks explaining short words. Another data set is TIMIT containing 6300 playlists of American English for phonetics. The most standard data set is VoxCeleb containing nearly 1,000,000 collections with 133Mb size describing human speech taken from different videos.

Video Data set

There are valuable data sets for video clippings. Some of them are TV Human Interaction Data set with 300 recordings and file size is 156MB of type MP4. The contents are various shows of social actions broadcasted in TV shows. Another data set is THUMOS with 25,000,000 collections with 385 KB size and the file type is MP4. The clippings consist of multi branded action classification. The exclusive data set is YouTube Face containing 3425 recordings designed for biometrics like face identification. The famous data set is 50 salads with 31 GB file size and the file type is RGB and the video collections are annotated data set of 25 participants engaged in food and beverage preparation. Above-described data sets are samples of the available data set for open access involved in event detection.

REPRESENTATIVES OF AUDIO AND VIDEO

The multimedia attributes like audio and video are used for automatic event detection in real life incidents. The properties that help in analysing and predicting the correct results are the representations like the features of audio and classification models used for predicting the actions with audio matching the video[69] for event identification. Some of the features are discussed in this section.

Audio features in Event detection

The sound received along with the video clippings or online videos are also engaged in detecting the events. The video actions are identified using sound [43-50] features also, in video films or any scene in the video. The features used for audio detection are numerous and compared as shown in Table 1.

Implementation of audio features

The features used for audio [51] detections are prescribed in the above table and the two important features are temporal which uses the signal waveform of audio for inquiry purpose. The other one is spectral method using the audio spectral demonstration process. The last feature is Prosodic which is used for audio signal awareness. For this type of feature any classifier can be used. [6]. Some of the features are discussed in brief They are as follows.

Zero Crossings

The zero crossing rates (ZCR) denote the frequency of the signal passing the zero signals per unit time. The frequency of the value of the signal changing from maximum to minimum divided by the length of the audio frame is the zero crossing. The equation is

$$ZC = \sum_{n=1}^{N} \frac{|sgnx(n) - sgn[x(n-1)]|}{2N}$$

Where sgn x(n) is the sign function.





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Mel-Frequency Cepstral Coefficients

This method uses a new scale called Mel which is calculated by a scale pitch similar in distance with each one. The signal bands wavelength is distributed based on the Mel scale which proves more accuracy. The cosine transform is calculated [52-60] to extract the coefficient from a single frame and the obtained sign is the Mel scale final signal. The equation is

 $C_n = \sum_{m=1}^{M} [\log x(m)] \cos[\pi n/m(m-\frac{1}{2})]$

Where final output is x(m) and m=1...M and n is the index of the coefficient.

Pitch

One of the prosodic features is the pitch providing useful aspects of audio like frequency, duration, loudness etc. Pitch is defined as the organisation of sinusoidal waves with basic intensity and amplitude. Audio is the combination of music, speech and various tones of sound containing pitch as the property. One type of pitch is absolute pitch which is the notes of the audio calculated by the number of pulsations per unit time. Pitch is used to estimate the peak of the amplitude to identify the particular tone.

Rhythm

Rhythm is also called duration provides characteristics of verbal and non -verbal audio identified by the nature of abstraction. They provide two types of attributes [7] namely the outlines of the sound fundamental notes and duration of each expression.

Harmonic Ratio (HR)

Another feature of audio used for event detection is the measure of the harmonic components within the range of continuum. It provides the high range of autocorrelation of the audio. These are some of the features of audio used for identification of events.

Classification models for audio and video processing

The next step after feature selection and extraction is the classification of the audio and video signals into different actions. The actions are characterised by the attributes of audio and video [70] to define the particular event automatically in the videos displayed. For this purpose, many classification models are built using bench mark data set. There are different categories of classification models and they are classified into three basic types namely unsupervised, supervised and semi supervised learning algorithms. This section deals with investigation of some classification models and their features.

Unsupervised Learning Approach

In this method the data set is untamed and not belongs to any class in prior. That is the data set is not labelled and the approach is to identify the class using some measures for detection. Using the similarity among the subject clustering is done to group similar data. So, cluster analysis is the fundamental attitude for unsupervised learning method. Some of the models are

Hierarchical and Partition clustering method

A clustering method used for making a large cluster or group by combining two or more subgroups. It can be both bottom to top or vice versa. Partition method deals with forming groups by changing the objects from one to another by dividing the group continuously. The number of clusters is decided earlier before partitioning.

Artificial Neural Network

The neural network system similar to human brain for classification using processors [62] and multiple connections inside the network. The most basic models are Self organizing map and adaptive resonance theory.





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Hidden Markov models

A graphical model which is based on probability theory to identify a group of similar objects those are unknown with the set of known variables. This is a statistical model used for classification based on internal factors for estimation.

Supervised Learning methods

In this method the data set objects belong to a particular class and are labelled for classification. This approach is best suited for classifying audio signals and there are many pre trained models used for classification. Some of the models are discussed in brief. They are

Artificial Neural Network

For audio and video [61] classification the weights of the network are finely tuned to find the hidden layer weights which are used for clarification. The popular models are multi-layer perceptron and Radial basis Function models used for classification

Instance based

The classification method based on the theory that samples can be used iteratively and one example is K nearest neighbour method. Here the values are predicted based on the neighbour around the value and Euclidean measure is used to calculate the similarity.

Random Forest

Random forest belongs to ensemble learning by building numerous trees from various samples and making decisions based on majority voting scheme for classification. It uses bagging and feature division method to predict any result based on the accumulation of all decision trees output.

Support vector machine

A classification method by adding data points to the feature space which are close to the hyper plane. It is the linear model to partition the data into classes based on the line of hyper plane.

Semi supervised learning methods

This hybrid method combines both unsupervised and supervised learning approaches for better accuracy. This method searches suitable classifier from both the procedures for classification. Some popular models used are

Self-training methods

A popular semi- supervised method trained with sample data which is labelled and used to test the unlabelled data. The main attributes are self- training with labelled data set and huge unlabelled events.

Co-Training and EM method

This method is suitable for small amount of known data with huge amount of unknown data. There are two classifiers for two different sets of data. Expectation and Maximization method also uses to cluster unknown data with already available values.

Transudative support machines

In recurrent neural network this method is applied for predicting problems by converting into another form.

Performance Evaluation

From the above given algorithms the best method is chosen by calculating the performance measures. There is basic four measures used. TPR is true positive rate for sensitivity gives actual positive values. TNR is true negative rate for





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specificity gives actual negative value. FPR for false positive rate is wrongly detecting true values and FNR is false negative rate wrongly predicting wrong values.

CONCLUSION

The final section of this paper is the conclusion part which summarises the survey of event detection methods using video and audio. This paper provided a general sketch of the need for automatic detection of events using audio and video with their common applications. Data set is discussed in the next section and the features used for classification is enumerated. The last section described the various models used for classification. A simple outline is discussed and this study can be extended by using any hybrid model for processing.

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Table 1. Categories of audio features

S.No	Feature Names	Sub Types
1	Temporal	Zero Crossings, Centroid, Roll off, Flux, Short term energy
2	Spectral	MFCC, Audio spectrum spread, centre, Flatness, Spectral entropy, signal bandwidth, Energy ratio, Linear prediction, MPEG7
3	Prosodic	Pitch, rhythm, intensity, Beat strength, Regularity of rhythm
4	Others	Harmonic ratio, root mean square, Time envelope, Low energy rate, Loudness





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RESEARCH ARTICLE

Bayesian Predictive Analysis for the Chris-Jerry Distribution with **Various Priors**

G.Meenakshi1 and B. Balachandar2*

¹Professor, Department of Statistics, Faculty of Annamalai University, Annamalai Nagar, Tamil Nadu,

²Research Scholar, Department of Statistics, Annamalai University, Annamalai Nagar, Tamil Nadu, India.

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*Address for Correspondence G Meenakshi

Professor, Department of Statistics,

Faculty of Annamalai University, Annamalai Nagar, Tamil Nadu, India.

Email: profgmau@gmail.com

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ABSTRACT

Bayesian predictive density is a crucial concept in Bayesian statistics, offering a probabilistic approach to prediction and decision-making under uncertainty. It incorporates existing knowledge and new data to create probability distributions over potential outcomes. This is particularly valuable when data is limited, as it provides a way to quantify uncertainty and update predictions over time. Bayesian predictive density aids in model selection, guiding the choice of the most fitting model, and it allows decision-makers to weigh potential risks and rewards comprehensively. Its flexibility accommodates prior beliefs and addresses heterogeneity in complex systems. By providing a range of possible outcomes along with their associated probabilities, it offers a practical tool for assessing and managing uncertainty. In sequential analyses, it ensures continuous adaptation to incoming data. Ultimately, Bayesian predictive density enables a deeper understanding of future events and supports informed decisions based on a holistic view of uncertain situations. this research concentrates on a new Bayesian predictive model for Chris-jerry distribution and its application in HIV viral replication.

Keywords: Chris-jerry distribution, posterior distribution, prior, predictive distribution.

INTRODUCTION

The landscape of Bayesian predictive models is a testament to the enduring power of mathematical inquiry and the evolution of probabilistic reasoning over centuries. From its humble beginnings in the 18th century, rooted in the pioneering contributions of Thomas Bayes and Pierre-Simon Laplace, Bayesian inference has undergone a





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remarkable transformation, shaped by mathematical development, philosophical debates, and the challenges posed by computational limitations and statistical paradigms. In the 20th century, despite its promising foundations, Bayesian predictive models faced significant hurdles. Computational constraints and the overwhelming influence of frequentist statistics hindered their widespread adoption, relegating them to the periphery of scientific inquiry. However, the tides began to turn with the advent of the Bayesian renaissance and the breakthroughs in computational techniques.

A pivotal moment in the resurgence of Bayesian predictive modeling came with the development of Markov Chain Monte Carlo (MCMC) methods, such as the Metropolis-Hastings algorithm and Gibbs sampler, in the 1950s and 1960s. These innovations provided a pathway to approximate complex posterior distributions, enabling the estimation of intricate Bayesian models and expanding their application horizon, particularly in the realm of prediction. The late 20th and early 21st centuries marked a watershed moment for Bayesian predictive modelling. Rapid advancements in computational capabilities, coupled with sophisticated algorithms, empowered researchers to implement highly intricate Bayesian models. This newfound computational prowess facilitated the derivation of predictive distributions in diverse fields, ranging from finance and machine learning to epidemiology, revolutionizing decision analysis by integrating uncertainty into critical decision-making processes. In this research the Bayesian predictive density and distribution is derived for Chris-Jerry distribution.

A Bayesian approach for Chris-jerry distribution

Chrisogonus K. Onyekwereand Okechukwu J. Obulezi 2022 have proposeda new one-parameter distribution named Chris-Jerry is suggested from a two-component mixture of Exponential (θ) distribution and Gamma $(3, \theta)$ distribution with mixing proportion $p = \theta / \theta + 2$ having a flexibility advantage in modeling lifetime data.

The probability distribution function of Chris-jerry distribution is given by

$$f(x) = \frac{\theta^2}{\theta + 2} \cdot (1 + \theta x^2) \cdot e^{-\theta x}$$

The likelihood function is given by

The likelihood function is given by
$$\pi(x_i) = \prod_{i=1}^n \frac{\theta^2}{\theta + 2} \cdot \left(1 + \theta x_i^2\right) \cdot e^{-\theta x_i}$$
$$= \frac{\theta^{2n}}{(\theta + 2)^n} (1 + \theta x_i)^n \cdot \left(e^{-\theta \sum_{i=1}^n x_i}\right)$$

Case: I

The prior is gamma prior (conjugate prior)

$$\mathbf{p}(\mathbf{\theta}) = \frac{\mathbf{e}^{-\mathbf{\theta}} \cdot \mathbf{\theta}^{\mathbf{r}-1}}{\mathbf{r}} \qquad \mathbf{r} > 0, \mathbf{\theta} > 0$$

The posterior distribution is given by
$$p\left(\frac{\theta}{x}\right) \propto \frac{\theta^{2n}}{(\theta+2)^n} (1+\theta x_i)^n . \left(e^{-\theta\sum_1^n x_i}\right) * \frac{e^{-\theta}.\theta^{r-1}}{\gamma r} \\ p\left(\frac{\theta}{x}\right) = \frac{1}{k} \frac{\theta^{2n}}{(\theta+2)^n} (1+\theta x_i)^n . \left(e^{-\theta\sum_1^n x_i}\right) * \frac{e^{-\theta}.\theta^{r-1}}{\gamma r}$$

$$\begin{split} k &= \int\limits_0^\infty \frac{\theta^{2n}}{(\theta+2)^n} (1+\theta x_i)^n. \left(e^{-\theta\sum_1^n x_i}\right) * \frac{e^{-\theta}.\theta^{r-1}}{\gamma r} d\theta \\ &= \frac{1}{2^n} \int\limits_0^\infty \frac{\theta^{2n}}{\left(1+\frac{\theta}{2}\right)^n} (1+\theta x_i)^n. \left(e^{-\theta\sum_1^n x_i}\right) * \frac{e^{-\theta}.\theta^{r-1}}{\gamma r} d\theta \\ &\left(1+\frac{\theta}{2}\right)^{-n} &= \sum_{i=0}^\infty (-1)^j c(n+j-1,j) \left(\frac{\theta}{2}\right)^j \end{split}$$





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$$\begin{split} (1+\theta x_i)^n &= \sum_{l=0}^\infty c(n,i)(\theta x)^l \\ &= \frac{1}{2^n} \int\limits_0^\infty \theta^{2n} \sum_{j=0}^\infty (-1)^j c(n+j-1,j) \left(\frac{\theta}{2}\right)^j \sum_{l=0}^\infty c(n,i)(\theta x)^l * \left(e^{-\theta \sum_1^n x_i}\right) * \frac{e^{-\theta} \cdot \theta^{r-1}}{\gamma r} d\theta \\ &= \sum_{j=0}^\infty (-1)^j c(n+j-1,j) \sum_{l=0}^\infty c(n,i)(x)^l \frac{1}{2^{n+j} [\gamma r]} \int\limits_0^\infty \theta^{2n+l+j+r-1} * \left(e^{-\theta (\sum_1^n x_i+1)}\right) d\theta \\ K &= \sum_{j=0}^\infty (-1)^j c(n+j-1,j) \sum_{l=0}^\infty c(n,i)(x)^l \frac{\{\gamma (2n+l+j+r)\}}{2^{n+j} [\gamma r] [\sum_1^n x_i + 1]^{2n+l+j+r}} \end{split}$$

The posterior distribution is given by

$$p\left(\frac{\theta}{x}\right) = \frac{1}{k} \frac{\theta^{2n}}{(\theta+2)^n} (1+\theta x_i)^n . \left(e^{-\theta \sum_1^n x_i}\right) * \frac{e^{-\theta} . \theta^{r-1}}{\gamma_r}$$

$$K = \sum_{i=0}^{\infty} (-1)^{j} c(n+j-1,j) \sum_{l=0}^{\infty} c(n,i)(x)^{l} \frac{\{\Upsilon(2n+l+j+r)\}}{2^{n+j} [\Upsilon r] [\sum_{1}^{n} x_{i}+1]^{2n+l+j+r}}$$

The predictive density is giv

$$\begin{split} g(x) &= \int\limits_{0}^{\infty} \prod_{i=1}^{n} \frac{\theta^{2}}{\theta+2} \cdot \left(1+\theta x_{i}^{2}\right) \cdot e^{-\theta x_{i}} * \frac{\theta^{2}}{\theta+2} \cdot \left(1+\theta x^{2}\right) \cdot e^{-\theta x} * \frac{e^{-\theta} \cdot \theta^{r-1}}{\gamma r} d\theta \\ &= \int\limits_{0}^{\infty} \prod_{i=1}^{n+1} \frac{\theta^{2}}{\theta+2} \cdot \left(1+\theta x_{i}^{2}\right) \cdot e^{-\theta x_{i}} * \frac{e^{-\theta} \cdot \theta^{r-1}}{\gamma r} d\theta \\ &= \int\limits_{0}^{\infty} \frac{\theta^{2(n+1)}}{(\theta+2)^{(n+1)}} (1+\theta x_{i})^{(n+1)} \cdot \left(e^{-\theta \sum_{1}^{n+1} x_{i}}\right) * \frac{e^{-\theta} \cdot \theta^{r-1}}{\gamma r} d\theta \end{split}$$

$$\begin{split} &= \int\limits_0^\infty \frac{\theta^{2(m)}}{(\theta+2)^{(m)}} (1+\theta x_i)^{(m)}. \left(e^{-\theta\sum_1^m x_i}\right) * \frac{e^{-\theta}.\,\theta^{r-1}}{\gamma r} d\theta \\ &= \sum\limits_{j=0}^\infty (-1)^j c(m+j-1,j) \sum\limits_{l=0}^\infty c(m,i)(x)^l \frac{1}{2^{n+j} [\gamma r]} \int\limits_0^\infty \theta^{2m+l+j+r-1} * \left(e^{-\theta(\sum_1^m x_i+1)}\right) d\theta \\ g(x) &= \sum\limits_{j=0}^\infty * \sum\limits_{l=0}^\infty \left\{c(m,i)(x)^l\right\} \!\! \left\{(-1)^j c(m+j-1,j)\right\} \!\! \left[\frac{\{\gamma(2m+l+j+r)\}}{2^{m+j} [\gamma r] [mx+1]^{2m+l+j+r}}\right] \end{split}$$

Case:II

Let the prior distribution is consider as quasi prior

$$p(\theta) = \frac{1}{\theta^d}$$

The joint posterior density is given by

The joint posterior density is given by
$$p(\theta/x) \propto \frac{1}{\theta^d} * \frac{\theta^{2n}}{(\theta+2)^n} (1+\theta x_i)^n . \left(e^{-\theta \sum_1^n x_i}\right)$$
$$= \frac{1}{K} \frac{1}{\theta^d} * \frac{\theta^{2n}}{(\theta+2)^n} (1+\theta x_i)^n . \left(e^{-\theta \sum_1^n x_i}\right)$$

$$K = \int\limits_0^\infty \frac{1}{\theta^d} * \frac{\theta^{2n}}{(\theta+2)^n} (1+\theta x_i)^n. \left(e^{-\theta \sum_1^n x_i}\right) d\theta$$

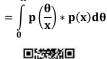




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$$\begin{split} & \text{Let} \\ & a = \sum_{i=0}^{n} x_{i} \\ & K = \int_{0}^{\infty} \frac{1}{\theta^{d}} * \frac{\theta^{2n}}{(\theta+2)^{n}} (1+\theta x_{i})^{n}. \left(e^{-\theta a}\right) d\theta \\ & \text{Where} \\ & (1+\theta x)^{n} = \sum_{i=0}^{\infty} c(n,i)(\theta x)^{i} \\ & K = \int_{0}^{\infty} \frac{1}{\theta^{d}} * \frac{\theta^{2n}}{(\theta+2)^{n}} * \sum_{i=0}^{\infty} c(n,i)(\theta x)^{i} * \left(e^{-\theta a}\right) d\theta \\ & \sum_{i=0}^{\infty} c(n,i) x^{i} = b \\ & K = b \int_{0}^{\infty} \frac{1}{\theta^{d}} * \frac{\theta^{2n}}{(\theta+2)^{n}} * \theta^{i} \left(e^{-\theta a}\right) d\theta \\ & = b \int_{0}^{\infty} \frac{\theta^{2n+i-d}}{(\theta+2)^{n}} * \left(e^{-\theta a}\right) * \left(\theta+2\right)^{-n} d\theta \\ & = b \int_{0}^{\infty} \theta^{2n+i-d} * \left(e^{-\theta a}\right) * \left(\theta+2\right)^{-n} d\theta \\ & = \frac{b}{2^{n}} \int_{0}^{\infty} \theta^{2n+i-d} * \left(e^{-\theta a}\right) * \sum_{j=0}^{\infty} (-1)^{j} \binom{n+j-1}{j} \left(\frac{\theta}{2}\right)^{j} d\theta \\ & \left(1+\frac{\theta}{2}\right)^{-n} = \sum_{j=0}^{\infty} (-1)^{j} \binom{n+j-1}{j} \left(\frac{\theta}{2}\right)^{j} d\theta \\ & \sum_{j=0}^{\infty} (-1)^{j} \binom{n+j-1}{j} \left(\frac{1}{2}\right)^{j} = e \\ & = \frac{b}{2^{n+j}} \int_{0}^{\infty} \theta^{2n+i-d} * \left(e^{-\theta a}\right) * \sum_{j=0}^{\infty} (-1)^{j} \binom{n+j-1}{j} \left(\theta\right)^{j} d\theta \\ & = \frac{b}{2^{n}} \int_{0}^{\infty} \theta^{2n+i-d+j} * \left(e^{-\theta a}\right) d\theta \\ & K = \frac{b}{2^{n}} \int_{0}^{\infty} \theta^{2n+i-d+j} * \left(e^{-\theta a}\right) d\theta \\ & K = \frac{b}{2^{n}} \int_{0}^{\infty} \theta^{2n+i-d+j} * \left(e^{-\theta a}\right) d\theta \\ & K = \frac{b}{2^{n}} \int_{0}^{\infty} \theta^{2n+i-d+j} * \left(e^{-\theta a}\right) d\theta \\ & K = \frac{b}{2^{n}} \int_{0}^{\infty} \theta^{2n+i-d+j} * \left(e^{-\theta a}\right) d\theta \\ & K = \frac{b}{2^{n}} \int_{0}^{\infty} \theta^{2n+i-d+j} * \left(e^{-\theta a}\right) d\theta \\ & K = \frac{b}{2^{n}} \int_{0}^{\infty} \theta^{2n+i-d+j} * \left(e^{-\theta a}\right) d\theta \\ & K = \frac{b}{2^{n}} \left(e^{-\theta a}\right) e^{2n+i-d+j} * \left(e^{-\theta a}\right) d\theta \\ & K = \frac{b}{2^{n}} \left(e^{-\theta a}\right) e^{2n+i-d+j} * \left(e^{-\theta a}\right) d\theta \\ & K = \frac{b}{2^{n}} \left(e^{-\theta a}\right) e^{2n+i-d+j} * \left(e^{-\theta a}\right) d\theta \\ & E = \frac{b}{2^{n}} \left(e^{-\theta a}\right) e^{2n+i-d+j} * \left(e^{-\theta a}\right) d\theta \\ & E = \frac{b}{2^{n}} \left(e^{2n+i-d+j} * \left(e^{-\theta a}\right) d\theta \right) d\theta \\ & E = \frac{b}{2^{n}} \left(e^{2n+i-d+j} * \left(e^{-\theta a}\right) d\theta \right) d\theta \\ & E = \frac{b}{2^{n}} \left(e^{2n+i-d+j} * \left(e^{-\theta a}\right) d\theta \right) d\theta \\ & E = \frac{b}{2^{n}} \left(e^{2n+i-d+j} * \left(e^{-\theta a}\right) d\theta \right) d\theta \\ & E = \frac{b}{2^{n}} \left(e^{2n+i-d+j} * \left(e^{-\theta a}\right) d\theta \right) d\theta \\ & E = \frac{b}{2^{n}} \left(e^{2n+i-d+j} * \left(e^{-\theta a}\right) d\theta \right) d\theta \\ & E = \frac{b}{2^{n}} \left(e^{2n+i-d+j} * \left(e^{2n+i-d+j} * \left(e^{-\theta a}\right) d\theta \right) d\theta \\ & E = \frac{b}{2^{n}} \left(e^{2n+i-d+j} * \left(e^{2n+i-d+j} *$$

The predictive density is given by







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$$\begin{split} &=\int_0^\infty \frac{1}{K} \frac{1}{\theta^4} * \frac{\theta^{2n}}{(\theta+2)^n} (1+\theta x_l)^n, \left(e^{-\theta \sum_l^n x_l}\right) * \frac{\theta^2}{\theta+2}, \left(1+\theta x^2\right), e^{-\theta x} d\theta \\ &=\int_0^\infty \frac{1}{K} \frac{1}{\theta^4} * \frac{\theta^{2n+1}}{(\theta+2)^{n+1}} (1+\theta x_l)^{n+1}, \left(e^{-\theta \sum_l^{n+1} x_l}\right) d\theta \\ &=\int_0^\infty \frac{1}{K} \frac{\theta^{2n+1-d}}{(\theta+2)^{n+1}} (1+\theta x_l)^{n+1}, \left(e^{-\theta \sum_l^{n+1} x_l}\right) d\theta \\ &=\frac{1}{K} \int_0^\infty \frac{\theta^{2n+1-d}}{(\theta+2)^{n+1}} (1+\theta x_l)^{n+1}, \left(e^{-\theta \sum_l^{n+1} x_l}\right) d\theta \\ &(1+\theta x_l)^{n+1} = \sum_{l=0}^\infty c(n+1,l)(\theta x)^l, \left(e^{-\theta \sum_l^{n+1} x_l}\right) d\theta \\ &=\frac{1}{K} \int_0^\infty \frac{\theta^{2n+1-d}}{(\theta+2)^{n+1}} * \theta^l * g * \left(e^{-\theta \sum_l^{n+1} x_l}\right) d\theta \\ &=\frac{1}{K} \int_0^\infty \frac{\theta^{2n+1-d+1}}{(\theta+2)^{n+1}} \left(e^{-\theta \sum_l^{n+1} x_l}\right) d\theta \\ &=\frac{g}{2^n K} \int_0^\infty \frac{\theta^{2n+1-d+1}}{(\theta+2)^{n+1}} \left(e^{-\theta \sum_l^{n+1} x_l}\right) d\theta \\ &=\frac{g}{2^n K} \int_0^\infty \frac{\theta^{2n+1-d+1}}{(\theta+2)^{n+1}} \left(e^{-\theta \sum_l^{n+1} x_l}\right) d\theta \\ &=\frac{g}{2^n K} \int_0^\infty \theta^{2n+1-d+1} * (1+\theta)^{-(n+1)} * \left(e^{-\theta \sum_l^{n+1} x_l}\right) d\theta \\ &=\frac{g}{2^n K} \int_0^\infty \theta^{2n+1-d+1} * (1+\theta)^{-(n+1)} * \left(e^{-\theta \sum_l^{n+1} x_l}\right) d\theta \\ &=\frac{g}{2^n K} \int_0^\infty \theta^{2n+1-d+1} * \sum_{m=0}^\infty (-1)^l \binom{n+m-2}{m} \left(\frac{\theta}{2}\right)^m * \left(e^{-\theta \sum_l^{n+1} x_l}\right) d\theta \\ &=\frac{g}{2^n K} \int_0^\infty \theta^{2n+1-d+1} * \left(e^{-\theta \sum_l^{n+1} x_l}\right) d\theta \\ &=\frac{g}{2^n K} \int_0^\infty \theta^{2n+1-d+1} * \left(e^{-\theta \sum_l^{n+1} x_l}\right) d\theta \\ &=\frac{g}{2^n K} \int_0^\infty \theta^{2n+1-d+1} * \left(e^{-\theta \sum_l^{n+1} x_l}\right) d\theta \\ &=\frac{g}{2^n K} \int_0^\infty \theta^{2n+1-d+1} * \left(e^{-\theta \sum_l^{n+1} x_l}\right) d\theta \\ &=\frac{g}{2^n K} \int_0^\infty \theta^{2n+1-d+1} * \left(e^{-\theta \sum_l^{n+1} x_l}\right) d\theta \\ &=\frac{g}{2^n K} \int_0^\infty \theta^{2n+1-d+1} * \left(e^{-\theta \sum_l^{n+1} x_l}\right) d\theta \\ &=\frac{g}{2^n K} \int_0^\infty \theta^{2n+1-d+1} * \left(e^{-\theta \sum_l^{n+1} x_l}\right) d\theta \\ &=\frac{g}{2^n K} \int_0^\infty \theta^{2n+1-d+1} * \left(e^{-\theta \sum_l^{n+1} x_l}\right) d\theta \\ &=\frac{g}{2^n K} \int_0^\infty \theta^{2n+1-d+1} * \left(e^{-\theta \sum_l^{n+1} x_l}\right) d\theta \\ &=\frac{g}{2^n K} \int_0^\infty \theta^{2n+1-d+1} * \left(e^{-\theta \sum_l^{n+1} x_l}\right) d\theta \\ &=\frac{g}{2^n K} \int_0^\infty \theta^{2n+1-d+1} * \left(e^{-\theta \sum_l^{n+1} x_l}\right) d\theta \\ &=\frac{g}{2^n K} \int_0^\infty \theta^{2n+1-d+1} * \left(e^{-\theta \sum_l^{n+1} x_l}\right) d\theta \\ &=\frac{g}{2^n K} \int_0^\infty \theta^{2n+1-d+1} * \left(e^{-\theta \sum_l^{n+1} x_l}\right) d\theta \\ &=\frac{g}{2^n K} \int_0^\infty \theta^{2n+1-d+1} * \left(e^{-\theta \sum_l^{n+1} x_l}\right) d\theta \\ &=\frac{g}{2^n K} \int_0^\infty \theta^{2n+1-d+1} * \left(e^{-\theta \sum_l^{n+1} x_l}\right) d$$

The predictive density with quasi prior is given by

$$g(\theta/x) = \frac{g * h}{2^{n}K(a)^{2n+d+l+m+2}} [\Gamma(2n+d+l+m+2)])$$





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The predictive density with diffuse prior is given by d=0

$$g(\theta/x) = \frac{g * h}{2^n K(a)^{2n+l+m+2}} [\Gamma(2n+l+m+2)])$$

The predictive density with non-infective prior is given by d=1

$$g(\theta/x) = \frac{g*h}{2^n \mathsf{K}(a)^{2n+l+m+3}} [\Gamma(2n+l+m+3)])$$

The predictive distribution is given by

$$\begin{split} &G(\theta/x) = \int_{0}^{x} \frac{\sum_{l=0}^{\infty} c(n+1,l)(x)^{l} * h}{2^{n} Ka} [\Gamma(2n+l+m+3)]) dx \\ &= \int_{0}^{x} \frac{\sum_{l=0}^{\infty} c(n+1,l)(x)^{l} * h}{(\sum_{i=0}^{\infty} c(n,i)x^{i} \cdot e)\Gamma(2n+i+j-d+1)} [\Gamma(2n+l+m+3)]) dx \\ &= \frac{\sum_{l=0}^{\infty} c(n+1,l)}{\Gamma(2n+i+j-d+1)e} \int_{0}^{x} \frac{(x)^{l}}{(\sum_{i=0}^{\infty} c(n,i)x^{i} \cdot)} [\Gamma(2n+l+m+3)]) dx \\ &= \frac{\sum_{l=0}^{\infty} c(n+1,l)}{\Gamma(2n+i+j-d+1)e} \int_{0}^{x} \frac{(x)^{l-i}}{(\sum_{i=0}^{\infty} c(n,i))} [\Gamma(2n+l+m+3)]) dx \\ &= \frac{\sum_{l=0}^{\infty} c(n+1,l)}{\Gamma(2n+i+j-d+1)e} \int_{0}^{x} \frac{(x)^{l-i}}{(\sum_{i=0}^{\infty} c(n,i))} [\Gamma(2n+l+m+3)]) dx \\ &= \frac{\sum_{l=0}^{\infty} c(n+1,l)}{\Gamma(2n+i+j-d+1)e} [\Gamma(2n+l+m+3)](x)^{l-i+1} \\ &= \frac{\sum_{l=0}^{\infty} c(n+1,l)}{\Gamma(2n+i+j-d+1)e} [\sum_{i=0}^{\infty} c(n,i))(l-i+1) \end{split}$$

Analysis

The above plots shows that the value predictive density decrease as value of the random variable (x) increases, specifically for the conjugate prior. Conversely, for quasi, non-informative, and diffuse priors, the predictive density decreases with an increase in the random variable

CONCLUSION

The depicted plots reveal a discernible pattern: as the random variable (x) increases, the predictive density experiences a noticeable decrease. This trend is particularly evident in the conjugate prior, presenting a distinct behavior compared to quasi, non-informative, and diffuse priors. For the latter group, the predictive density also decreases with an ascending random variable, albeit with differences in the specific patterns. These findings underscore the substantial impact of prior selection on the predictive dynamics of the Chris-Jerry distribution. The distinct trend in predictive density observed in the conjugate prior highlights the sensitivity of the model to different prior specifications. In conclusion, the careful selection of a prior for the Chris-Jerry distribution proves essential for precise probability predictions concerning the event of interest. The variations in predictive density emphasize the nuanced role of prior choice in influencing the model's performance. This insight is invaluable for practitioners grappling with the intricacies of the Chris-Jerry distribution in diverse applications. Ultimately, this research serves as a guide for practitioners, emphasizing the critical role of prior selection in optimizing the accuracy of predictions within the Chris-Jerry distribution framework. The observed variations provide actionable insights, empowering decision-makers to make informed choices in real-world scenarios.





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Table 1: Predictive density

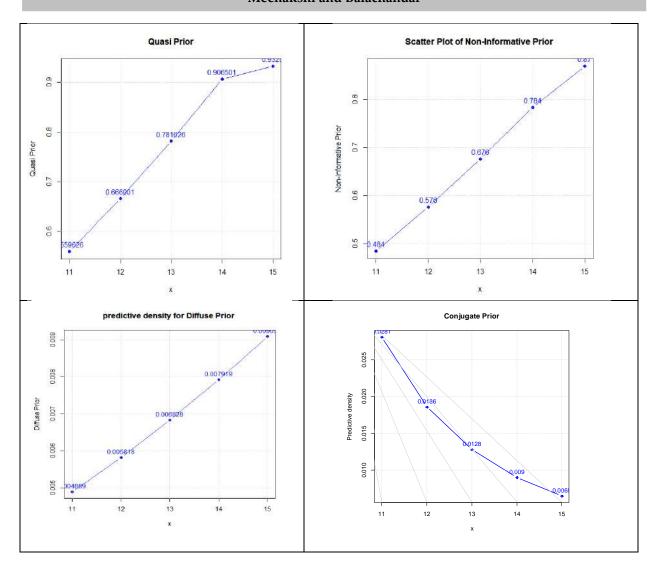
X	Quasi prior	Non-informative prior	diffuse prior	Conjugate prior
11	5.60E-01	0.484	0.004889	0.0281
12	6.66E-01	0.576	0.005818	0.0186
13	7.82E-01	0.676	0.006828	0.0128
14	9.07E-01	0.784	0.007919	0.0090035
15	9.11E-01	0.87	0.009091	0.0064779





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RESEARCH ARTICLE

Nutritional and Functional Evaluation of Pomegranate Peel Powder **Incorporated Waffles**

S. Mohana¹ and Arivuchudar.R^{2*}

¹Student, Department of Nutrition and Dietetics, Periyar University, Salem, Tamil Nadu, India. ²Assistant Professor, Department of Nutrition and Dietetics, Periyar University, Salem, Tamil Nadu, India.

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*Address for Correspondence Arivuchudar.R

Assistant Professor, Department of Nutrition and Dietetics, Periyar University, Salem, Tamil Nadu, India. Email: achudar24@gmail.com



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ABSTRACT

Sustainability entails addressing the demands without jeopardising future generation's ability to meet their needs. It is more than just being environmentally conscious. Most conceptions of sustainability include considerations for social equality and economic development. Sustainable food management is a systematic method that tries to reduce food waste and its associated implications throughout the life cycle. Pomegranate cultivation and consumption have always increased due to their flavour and nutrients. A huge number of byproducts are formed during fruit processing, such as peels and seeds which can cause environmental contamination if not handled appropriately. The study was carried out to formulate value-added waffles, by incorporating pomegranate peel powder (Punica granatum) in different proportions viz. V1 (90g wheat flour + 10g pomegranate peel powder) and V2 (85g wheat flour + 15 g pomegranate peel powder), V3 (80g wheat flour+ 20g pomegranate peel powder) respectively for each variation of the product. On the basis of sensory evaluation carried out using the nine-point hedonic scale, it was observed that the variation V2 scored best regarding appearance, colour, texture, taste, flavour and overall acceptability. Hence, the accepted variation V2 was subjected to nutritional and phytochemical analysis. The waffle was found to be higher in terms of nutritional parameters when compared with control and was also found to be a source of phytochemicals such as total flavonoids and phenols. Tannins were present in permissible limits, contributing to their extensive functions like antiinflammatory, anti-cancer, anti-bacterial and cardiovascular protection.

Keywords: Environmental sustainability, Food waste, Pomegranate peel, Waffles





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INTRODUCTION

The year 2021 was observed as the International Year of Fruits and Vegetables by the United Nations Organisation to propagate the nutritional importance of fruits and vegetables for healthy living. Of late, now it is understood that not only the whole fruits and vegetables, but the by-products from the fruits and vegetable markets and processing industries serve as a treasure of nutrients. Numerous studies state that the peels that are discarded as waste pose a threat to the environment, rather they can be used as value enhancers in different food formulations and drug formulations. To mention, a few studies, *Ananas cosmosus* fruit peel has anti-inflammatory activity [1], lemon peel extract can be used as a natural binder in aspirin tablets [2], custard peel extract and carica papaya peel extract exhibit antimicrobial and antioxidant property [3,4], pomegranate peels were used to develop phytotomes, for novel drug delivery system [5] and as a cosmetic ingredient in herbal lipsticks [6].Pomegranate peel finds application as ointment base for burn, wound and scar healing [7]. The pomegranate being a colorful fruit, grabs attention. Pomegranate is the fruit of a shrub (*Punica granatum. L*), grown predominantly in west Asia and the Mediterranean region, as well as other parts of the world, including America, where the environment is favorable for its growth [8].

Pomegranate fruits have a variety of constituents in their seeds, skins, and arils, which play a therapeutic role in the management of health through the control of numerous biological processes. Over 60% of the pomegranate's weight is made up of its peel, and is a rich source of fibre, minerals like calcium, magnesium, phosphorus, potassium, and sodium as well as a variety of phytochemicals like ellagitannins, proanthocyanidin compounds, and flavonoids [9]. Pomegranate waste is generated at all stages of the fruit's life cycle, including agricultural production, industrial manufacturing, and processing. Pomegranate by-products can be used since they are high in bioactive substances. Pomegranate peel being a significant by-product of pomegranate juice production, the presence of polyphenols, flavonoids, and hydrolysed tannins are responsible for its varied biological functions [10]. Pomegranate peel attracts attention due to its wound healing capacity, as peels contain high amounts of bioactive compounds which inhibit migration of Salmonella on wet surfaces. Pomegranate peels are also characterized by the presence of gallic acid, ellagic acid and punicalagin. They possess free radical scavenging properties and antibacterial activities against enteric pathogens like *E.coli*, *Salmonella species*, *Shigella species*, *Vibrio cholera*, viruses, fungi, and mould [11].

Pomegranate peel has a great therapeutic effect on chronic inflammation, like ulcerative colitis, blood pressure reduction, anti-diabetes, and anticancer [12]. Literature states that both the fresh and dried pomegranate peel powder is enriched with macro and micro minerals like calcium, potassium, phosphorus, sodium, iron, zinc, manganese, selenium and copper [13]. Pomegranate peel can be used as a functional ingredient because it is a good source of crude fibres, which have numerous health benefits such as the ability to lower serum LDL-cholesterol levels, improve glucose tolerance and insulin response, reduce hyperlipidaemia and hypertension, contribute to gastrointestinal health, and prevent certain cancers such as colon cancer [14]. The invitro and invivo studies using the pomegranate peel extract had shown to exhibit potent activity against helminthic, diarrhea, tumor growth and was immuno-stimulatory and hepato-Protective in nature [15]. Anti diabetic activity of pomegranate peel was studied in diabetic rats and the effect was comparable with oral hypoglycemic drugs [16]. Pectin, a valuable food additive can also be extracted from pomegranate peel [17]. Taking into consideration, the nutritional richness of pomegranate peel powder, it was proposed in this study, to formulate value-added waffles, the most preferred young-generation snack product, by incorporating pomegranate peel powder.

MATERIALS AND METHODS

Procurement and processing of raw materials

The ingredients required for the formulation of pomegranate peel powder incorporated value added waffles, like whole wheat flour, pomegranate, coconut oil, baking powder, baking soda, sugar were procured from the local market in Salem, Tamil Nadu.





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Processing of pomegranate peel powder

Pomegranates were purchased and checked for any infestation or damage present. Pomegranate which was free from damage was peeled and then sun dried for 2-3 days, and ground into powder.

Formulation of pomegranate peel powder incorporated waffles

Value added batter was prepared using the ingredients listed in Table 1. Waffles was formulated as shown in table 1, for each variation, viz. Control, Variation 1, Variation 2, and Variation 3.

Acceptability of the product by organoleptic evaluation

The developed variations of the waffles were subjected to organoleptic evaluation to assess the maximum acceptability by the panel members. The quality attributes in terms of colour, appearance, flavour, texture, taste and overall acceptability were evaluated by semi-trained judges using score card with 9-point hedonic scale.

Nutritional and Phytochemical analysis of the accepted variation of pomegranate peel powder incorporated waffles

The nutritive value of accepted variation of waffle was assessed. The nutritional analysis of waffles was done using NSI diet calculator and standard laboratory procedures. The nutrients assessed include energy, carbohydrate, protein, fat, dietary fiber, iron, calcium, potassium, zinc, folic acid and total sugar. The phytochemicals like phenolic compounds, flavonoids and alkaloids were also assessed.

RESULTS AND DISCUSSION

The results of the formulated pomegranate peel powder incorporated waffles is presented and discussed further.

Organoleptic evaluation of the developed variations of the pomegranate peel powder incorporated waffles

Results of ANOVA Duncan multiple range tests showed that there was a significant difference ($p\le0.05$) between the control and all the variations of the waffles on the basis of appearance, color, texture, taste, flavor and overall acceptability. Results of sensory evaluation of the waffles prepared with 10%, 15%, and 20% pomegranate peel powder incorporation and control waffles are presented in table 2.

Appearance

The mean and standard deviation of control and developed variations was calculated with the sensory evaluation score given by panel members. 'Appearance' of the developed variations of waffles was analyzed according to its size, shape, and visual appearance, during sensory evaluation. Variation 2 got maximum score of 8.50±0.57,thus it was highly accepted by the panel members.

Color

Color is the most important sensory property of food product. Color gives the impression about the freshness, flavor and quality of the product. The mean and standard deviation of control and developed variations was calculated with the sensory evaluation score given by panel members. 'Color' of the developed variations was analyzed according to its color and visual appearance, in the sensory evaluation Variation 2 got maximum score of 8.17±1.08, hence it was highly accepted by the panel members. In a similar rice berry incorporated waffles study, malitol and palm sugar were substituted for sugar and no significant difference was observed in terms of appearance and colour [18].

Texture

Texture refers to those qualities of a food that can be felt with the fingers, tongue, palate or teeth. The waffle's texture is sensed by hardness and fracture ability. From the developed variations of waffles, variation 2 attained its





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maximum fracture ability, with the score 7.97 ± 0.81 , which is nearly equal to the score 8.27 ± 0.87 obtained by control. Hence variation 2 was highly accepted by the panel members.

Taste

The taste of food is caused by its chemical compounds. These compounds interact with the sensory (receptors) cells in the taste buds and can different types of tastes such as sweet, sour, salty, bitter, and savory. From the developed variations of waffles, variation 2 attained the taste of waffles, with score of 7.40±1.00, which is nearly equal to the score 7.93±0.78 obtained by control. Hence variation 2 was highly accepted by the panel members.

Flavor

Flavor of the particular food is also determined by the aromas picked up by nose. The primary function of the flavor is to add taste to the food. From the developed variations of waffles, variation 2 attained the flavors of waffles, with the score of 8.07±0.78, which is nearly equal to the score 8.50±0.57 obtained by control. Hence variation 2 was highly acceptable when compared to control. The incorporation of almond skin imparted a roasted flavor to waffles even without actual roasting and had significant impact in the flavor profile assessment by the panel members [19].

Overall acceptability

Overall acceptability of the food product is an important factor that is influenced by the sensory quality of the product. In the sensory evaluation, variation 2 got maximum score in comparison with control, V1 and V3. Variation 2 scores were observed to be highest of all the sensory characteristics with an overall acceptability of 8.13±.82, representing that variation 2 was highly accepted by the panel members.

Nutritional analysis of the accepted variation of pomegranate peel powder incorporated waffles

The above table infers waffles prepared with pomegranate peel powder is higher in energy content than control which is made from refined wheat flour, the protein content present in the accepted variation was 6.90g/100g which is greater than control of 218g/100g with the difference of +0.98g, the fat content is +7.08g greater in accepted variation than control 13.2g, the carbohydrate content was higher in accepted variation than control with 15.55g higher, the fiber was completely void in control when compared to the accepted variation with 1.44g, moisture content of accepted variation of waffle was 31.50g which higher than control. The total sugar content present in control was higher than accepted variation. The iron content of waffles prepared from pomegranate peel powder was found to be 41.42g/100g which was higher than control made from refined wheat flour with 1.7g/100g. The zinc content present in the waffles was higher in accepted variation (variation 2) with 9.89g/100g.when compared with control the sodium and potassium content was higher in accepted variation of waffle. The calcium content of waffles prepared from pomegranate peel powder was found to be 83.50g/100g which was lower than control. Similarly, millet flour incorporated waffle showed significant increase in total dietary fibre and minerals, and reduced carbohydrate content [20].

Phytochemical analysis of the accepted variation of the accepted variation of pomegranate peel powder incorporated waffles

The above table shows that the accepted variation of pomegranate peel powder incorporated waffles contains polyphenols, flavonoids, tannins, and alkaloids. It was also observed that raspberry incorporation increased phenolic compounds in waffles [21].

CONCLUSION

Pomegranate is the well-known fruit of a shrub (*Punica granatum*. *L*). It is known that the polyphenolic compounds of pomegranate peel have the most pronounced therapeutic effect. As nutritional and bioactive properties of this Pomegranate peel powder are better understood, the interest of these food is increasing. Peel powder has several health benefits including antioxidant properties so it is recommended for the people suffering from non-





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communicable disease. Waffles are usually easy to make and convenient to have around to pop in lunch boxes or for breakfast or as evening snacks. Hence pomegranate peel powder incorporated waffles will be suitable for people of all age groups. As pomegranate peels have demonstrated their potential as promising bioactive antioxidants, they may be a potential source of functional food ingredients and nutraceuticals that can be used in various areas of food industries.

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Table 1: Ingredients for the formulation of pomegranate peel powder waffles

	Level of incorporation			
Ingredients	Control	Variation 1	Variation 2	Variation 3
Whole wheat flour (g)	100	90	85	80
Pomegranate peel powder (g)	-	10	15	20
Sugar (g)	30	10	10	10
Milk (ml)	10	10	10	10
Vanilla extract (tsp)	2	1/2	1/2	1/2
Salt	A Pinch	A Pinch	A Pinch	A Pinch
Coconut Oil (ml)	3	3	3	3

Table 2: Statistical analysis of mean organoleptic values of developed waffles

j 0 1		<u>.</u>				
Variations	Appearance	Colour	Texture	Taste	Flavour	Overall acceptability
Control	8.03±0.67°	7.67±0.96 bc	8.27±0.87°	7.93±0.78 ^d	8.50±0.57 ^d	8.53±0.68 ^d
Variation 1	6.90±0.99ª	6.93±0.87ª	7.40±1.10 ^b	6.93±0.78 ^ь	7.33±0.76 ь	7.17±0.79 b
Variation 2	8.50±0.57 ^d	8.17±1.08°	7.97±0.81°	7.40±1.00°	8.07±0.78°	8.13±0.82°
Variation 3	7.50±0.97 b	7.27±1.08ab	6.83±0.95ª	6.33±0.96ª	6.57±0.93ª	6.77±0.77ª
Sig*	0.000 *	0.000 *	0.000 *	0.000 *	0.000 *	0.000 *

^{*} Significance at 5 % level. Values are expressed as mean \pm standard deviation. Samples with different superscripts within a column are significantly different from one another at (p \le 0.05).

Table 3: Comparison of nutrients between control and accepted variation of waffles (100 gms)

S. No	Parameters	Control	Accepted Variation (V2)	Difference
1	Energy(Kcal)	218	370.95	+152.95
2	Protein(g)	5.92	6.90	+0.98
3	Fat(g)	13.2	20.28	+7.08
4	Carbohydrate(g)	24.68	40.23	+15.55
5	Fiber content (g)	0	1.44	+1.44
6	Moisture (%)	31.50	30.83	-0.67
7	Total sugar (g)	24.68	18.34	-6.26



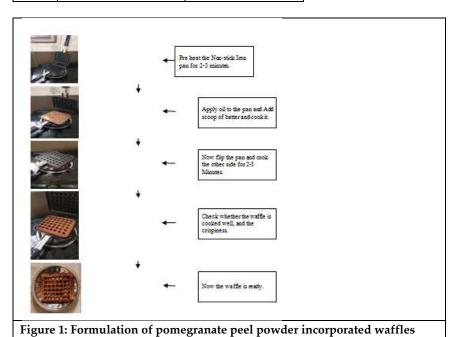


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8	Iron (mg)	1.7	41.42	+39.72
9	Zinc (mg)	0.51	9.89	+9.38
10	Potassium (mg)	119	1478.27	+1359.27
11	Calcium (mg)	191.25	83.50	-107.73

Table 4: Qualitative analysis of phytochemicals in accepted variation of waffles

S.NO	Qualitative Analysis	Present/ absent
1	Polyphenols	++
2	Flavonoids	++
3	Tannins	++
4	Alkaloids	++







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RESEARCH ARTICLE

Impact of Artificial Intelligence and Their Influence on Future **Workforces and Employment**

R. Prema^{1*} and K. Subramaniam²

¹Associate Professor, Department of Commerce, KPR College of Arts Science and Research (Affiliated to Bharathiar University) Coimbatore, Tamil Nadu, India.

²Associate Professor, Department of Commerce and Management, Kamban College of Arts and Science (Affiliated to Bharathiar University, Coimbatore) Tirupur, Tamil Nadu, India.

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*Address for Correspondence

R. Prema

Associate Professor, Department of Commerce, KPR College of Arts Science and Research (Affiliated to Bharathiar University), Coimbatore, Tamil Nadu, India. Email: rprems6@gmail.com & prema.r@kprcas.ac.in



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ABSTRACT

The rise of artificial intelligence (AI) has the potential to drastically impact the workforce in the near future. This paper explores the potential impact of AI onemployment and the future workforce, including both the positive and negative effects. On the positive side, AI can enhance productivity and efficiency, leading to increased economic growth and job creation. With the development of AI, businesses can automate repetitive tasks, freeing up employees to focus on more complex and creative work. In addition, AI can help companies make more informed decisions, leading to improved customer satisfaction and increased profits. However, there are also negative impacts of AI on employment. As AI technologies improve, they have the potential to replace human workers in certain industries. For example, manufacturing jobs have already been impacted by automation, and service jobs such as those in the transportation industry could be next. This could lead to a significant loss of jobs for low-skilled workers, who may struggle to find alternative employment. In addition, AI could exacerbate income inequality, as high-skilled workers who are able to work with and develop AI technologies may see their salaries rise while low-skilled workers are left behind. If AI algorithms are developed by a homogenous group of individuals, they may be biased against certain groups of people, leading to discrimination in hiring and other employment decisions. Additionally, as AI becomes more prevalent in the workforce, workers may feel a loss of control over their jobs and the decision-making process. This could lead to increased stress and burnout, as well as a decrease in job satisfaction. To mitigate the potential negative impacts of AI on employment, policymakers and businesses must take proactive steps. This includes investing in education and training programs to help workers develop the skills necessary to work alongside AI technologies, as well as developing regulations to prevent bias in AI algorithms.





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Additionally, businesses should be transparent about their use of AI and involve workers in the decision-making process to ensure that they feel valued and engaged. In conclusion, while AI has the potential to revolutionize the workforce and bring about positive change, it also has the potential to negatively impact employment and exacerbate income inequality. By taking proactive steps to mitigate the potential negative impacts, AI should be used in a responsible and equitable manner.

Keywords: AI in Employment, AI with Human Intelligence, Future Workforce and Intervention of AI.

INTRODUCTION

Every new discovery will have both benefits and drawbacks, but it is our responsibility as humans to manage this and make the most of the benefits to advance humankind and the globe. Artificial intelligence has great potential advantages. In order to keep the "rise of the robots" from getting out of control, humans must play a significant role. Others contend that artificial intelligence has the potential to wipe outhuman civilization if it gets in to the wrong hands. However, none of the AI programmes developed at that level had the capability to kill or enslave people. With autonomous cars and voice automation in homes, artificial intelligence has fast developed and is no longer just a concept from science fiction films and books. The predictions made for 2054 in the critically acclaimed film "Minority Report" are coming true more quickly than anticipated. In a study undertaken by experts at the University of Oxford, it was shown that artificial intelligence will exceed humans in a number of jobs by the years 2024, 2026, 2031, 2049, 2053, and 2031, including generating bestselling books, writing essays for school, and selling things. Artificial intelligence (AI) will inevitably outperform human intelligence in the coming years, affecting every area of our lives.

By employing facial recognition software rather than a key to access a hotel room, the face will function as the customer's identity, simplifying and streamlining daily interactions. Within minutes of placing an order, smaller drones will deliver therequested products right to our door. Virtual assistants powered by artificial intelligence will conduct human-like phone calls to set up an appointment. The time is not far away to have surgery performed by a robot. In a few years, a robot will do the procedure in place of a physical surgeon and help patients better comprehend their treatment options. These are only a few examples of the future-changing effects of artificial intelligence. Artificial intelligence future technology advancements may seem far off, yet they will come faster than anticipated. Leading tech firms are competing with one another to incorporate artificial intelligence into their everyday lives, which will open the door for a genuinely fantastic and exciting artificial intelligence future.

REVIEW OF LITERATURE

Moradi P. & Levy K. (2020) explored that many people have raised concern about the mass displacement of human workers as AI-driven technologies are being integrated into workplaces and labor procedures. The study offers a more nuanced perspective on the widely held belief that robots will replace humans in the workforce. We argue that economic predictions of significant job losses brought on by AI are of limited practical value because they frequently concentrate only on the technical aspects of task execution, ignoring broader contextual inquiry about the social aspects of work, organizational structures, and cross-industry effects. Shaukat K.(2020) examined the traits and behaviors of numerous robot types. The study examined that how robotics and humans are evolving together. By creating a working system that solves issues and produces good results and there is a need to better understand how the human brain works. Artificial intelligence is a huge field that is also making headway in the areas of business, healthcare, and quality control. Rauf, Muhammad Abdul (2021) instigates that as robots have made everyone's lives more laid-back and pleasant, there is an increasing amount of commonplace human-robot interaction. There is a need to understand how people and robots are evolving together. A number of field's and components have been included in this analysis as well as work done by several technicians and scientists. By





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developing an effective system that solves issues and produces positive outcomes, it is essential to have a better understanding how the human brain functions.

Boyd, Ross. (2021) discussed about the impact of AI-powered technologies on the 2016 US Presidential elections which were dominated by allegations of Russian meddling via large-scale troll farms, the leak of documents and emails from different Democratic Party servers, and the scandal involving Facebook and Cambridge Analytica. Throughout the 19th century, the work of significant political and economic philosophers vividly reflected the ambivalence that accompanied the early experiences of industrialization. In his 1930 essay, Keynes suggested that eliminating Adam's Curse will lead to the emergence of two more issues. Numerous experts have noted the issues with abstraction and lack of depth that come with using employment aggregates to gauge how sophisticated technology is affecting the workforce. Anagnoste, Sorin, (2021) stated that the companies are transferring their workloads to software robots through the use of automation, whether it be in the cloud, on- premise, or hybrid like. This transforms the outdated operating models, reduces labor, and creates employment that is better suited for people. With the rise in customer demand for self-services and the increasing dominance of messaging platforms in this environment, businesses are now able to create internal and external communications without the need for a new user interface. Modern chat bots are excellent enablers of human-to-robot contact, and their industry is anticipated to grow significantly over thecoming years. Chat bots can overcome difficulty and confusion in language thanks to the technologies that support natural language processing (NLP) and intent recognition. Ekaningrum, Nurul Efri, Muhammad Syahrul Hidayat, and Farida Yuliati. (2023) explained that nonetheless, this may jeopardize people's employment chances. This study was done to determine how automation and AI are affecting human jobs. The information used in this study was derived from a number of research findings and earlier investigations that continue to address the usage of AI and automation in the workplace. According to this study, automation and AI are currently displacing many employments. Yet, other aspects of human intelligence, like intuition and empathy, are still challenging for AI to replicate

Objectives of the Study

- To analyze the scope of AI in generating the employments.
- To know the impact of artificial intelligence and their influence on future workforces and employment.

RESEARCH METHOOLOGY

A descriptive research design was adopted for the study. Simple random sampling technique has been used for the study. The sample size considered for the study is 150. The area of the study is Tirupur and Coimbatore city. Tirupur is also known as the knitwear capital of India, accounting for 90% of India's cotton knitwear export. It contributes to a huge amount of foreign exchange in India. Coimbatore is the third largest city in the state and one of the most industrialized cities in Tamil Nadu, known as the textile capital of South India or the Manchester of the South India. The study includes quantitative data gathering, including questionnaires that gathered numerical data from a variety of people, including workers, housewives, businessmen, retired people, professional etc., These persons were deemed as sample for the study. For the study, both primary data and secondary has been taken into consideration. Primary data was gathered by using questionnaire and collected through Microsoft forms. Secondary data have been collected from different sources like journal, websites, magazines etc.

AI's Interference in Modern Period

Artificial intelligence, or AI, seems to be a popular topic right now. It has been noticed that job applicants are increasingly looking for AI expertise. In its simplest form, artificial intelligence combines computer science and huge data sets to facilitate problem-solving. Additionally, it incorporates the artificial intelligence subfields of deep learning and machine learning, which are frequently studied in tandem. These industries create expert systems that classify information or make predictions based on incoming data using AI algorithms. Without external input, the AI uses the facts presented to make decisions and choices that are appropriate to the environment. By and large, "AI"





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conjures up nightmares or science fiction fancies of computers conquering society. Although no one can predict with confidence how artificial intelligence (AI) will grow in the future, the current trends and advancements offer a very different picture of how AI will effect our daily lives. Artificial intelligence (AI) has been portrayed in the media in a broad variety of ways. In reality, artificial intelligence (AI) is already widely used and has an impact on a variety of things, including our search engine results to make purchases. Data indicates that during the past four years, the use of AI has grown by 270% in a variety of corporate industries.

A few examples of AI applications include advanced web search engines like google search, recommendation systems, voice assistants that can understand human speech like Siri and Alexa, self-driving cars like Waymo, generative or creative tools, automated decision making, and dominating the best strategy game systems (like chessandgo). Computers grow more skilled; tasks that were once considered to require intelligence are now routinely removed from the concept of AI. Rapid optical character recognition is frequently left out of definitions of AI, despite the fact that it is a widely used technique. There are several subfields in AI research, each with its own set of goals and resources. Natural language processing, planning, learning, reasoning, knowledge representation, vision, and the capacity to move and manipulate objects are among the traditional goals of AI research. General intelligence is one of the long-term goals in the area. Researchers in AI have used a variety of tools to approach this problem, including artificial neutral networks, formal logic, search and mathematical optimization, method-based statistics, probability, and economics. A wide range of other fields, such as computer science, psychology, linguistics, and philosophy, are also incorporated into AI. Artificial intelligence has seen numerous waves of hope, disappointment, and funding loss (AI winter), followed by new methods, successes, and increased funding since it was first established as an area of research in 1956. Since its inception, researchers have tried and failed with a wide range of approaches, including brain simulation, human problem-solving modeling, formal logic, a significant amount of information, and animal behavior imitation. In the first decade of the twenty-first century, very mathematical statistical machine learning has dominated the field, and this approach has been very successful in assisting with the resolution of many challenging problems in both industry and academia. Ancient Greece is where the concept of a "machine that thinks" first appeared however, significant developments and turning points in the development of artificial intelligence have occurred since the invention of electronic computing.

AI On Par Comparison with Human Intelligence

This field was founded on the theory that human intelligence may be so precisely described that a machine can be built to stimulate it. Since antiquity, myth, literature, and philosophy have all explored these issues. This sparked philosophical debates on the nature of the mind and the moral implications of creating artificial beings with human-like intelligence. Since then, computer scientists and philosophers have suggested that artificial intelligence (AI) may wind up becoming an extension of humans if it is not directed in using its logical abilities for the benefit of society. While traditional human activities are being displaced by digital life, human skills are increasing. Code-driven systems now have access to more than half of the world's population, opening up opportunities and threats that weren't previously imaginable. This is made possible by ambient information and connection.

The scientists predict that networked artificial intelligence would improve human performance but also endangering their autonomy, agency, and abilities. They talked about a variety of scenarios, including the possibility that computers could eventually match or even surpass human intelligence and capabilities in a variety of domains, including visual acuity, speech recognition, language translation, and sophisticated analytics. Examples of these domains include complex decision-making, reasoning, and learning. Smart systems would lower costs, save lives, and provide more custom-made features in automobiles, buildings and utilities, farms, and commercial operations. A number of the upbeat remarks focused on health care and the various ways Alcould be used to diagnose and cure patients, or to help seniors live longer, healthier lives. All could contribute to extensive public health initiatives based on the massive amounts of data that may be obtained in the approaching years regarding everything from nutrition to individual genomes. Additionally, some of the experts predicted that All would enable long-needed advancements in formal and informal education systems. However, whether they are optimistic or pessimistic, the majority of experts expressed concerns about how these new technologies may impact the core





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characteristics of what it means to be a human in the long run. In this non-scientific interview, the respondents were asked to express a precise reason as to why they believed AI would benefit or harm humanity.

AI's Current Position in the World

Artificial intelligence (AI) technologies are currently used in a wide range of industries, from consumer electronics to customer service software and technological arms races. Both the collection of patient data and the formulation of patient-specific health findings involve their utilization in the healthcare sector. During the COVID-19 Pandemic, AI techniques were utilized to monitor people's health behavior, which has the potential to save lives. Even in space, AI is assisting astronauts and looking for NASA scientists around the cosmos. It is employed in the creation of autonomous vehicles, in trade and banking, and even in military applications like as training and unmanned weaponry. The boundaries of AI applications are still being explored, and businesses, organizations, and institutions all around the world are integrating AI technology into their operations and advancements. The existing uses of AI have led to unfairness and ethical concerns, quite apart from the exciting insight into future possibilities they provide.

DISCUSSIONS AND FINDINGS

From the descriptive analysis, the researcher found that majority of the respondents are female, qualified with under graduation and they are in the age group of 20 to 30 years with the family annual income of between Rs 1,00,000 to 3,00,000. Majority of the respondents came to know about AI tools through social media in the past 2 to 3 years. They think future of AI will be exceptional and there will be more possibilities that AI will make humankind shudder. Most of the people think that our job will be secured in the future with AI and they opined that the intervention of AI in bank is good as the provided data cannot be tampered. They agreed with the AI's contribution in medical field without emotional conscience and have said that the human creativity cannot be achieved by AI. Majority of the respondents thought that creative, emotion-based activities cannot be replaced by AI, and the effect of AI in ten years will cause massive unemployment and AI will be beneficial to human race. They may choose AI along with external factors to make proper decision and any fraud and extra chargecan be avoided. By using analysis of variance, it is inferred that there is no significant difference in the mean value of revolution due to intervention of AI with age of the respondents. Majority of the respondents agreed that data-based works will be overtaken by AI. Further, the respondents opined that the prime problem faced due to the advancement of AI is in marketing functions and the benefit of AI is reduction in human error. Majority of the respondents thought that AI will gain success within '5 to 10 years'. It is inferred that majority of the respondents opined that the Customer service executive role will be greatly influenced by the AI. To find out the significant association between the educational qualification and the problems due to intervention of AI, chi-square test is applied. It has been found from Chi-Square test that p value (0.111) is more than 0.05, therefore the null hypothesis is accepted. Therefore, it is concluded that there is no significant association between the educational qualification and the problems due to intervention of AI.

To find out the impact of artificial intelligence on future workforce and employment, the multiple regression method is applied. In this, the impact of AI is the dependent variable, and its effect on Employment, the banking Sector, and Transport Sector are the independent variables. The above table depicts that multiple correlation co-efficients (R) indicate the impact of AI on future workforce and employment which will influence to affect the various sectors like employment, the banking sector and transport sector. The Co-efficients of determination (R²) indicates 22.7% impact on artificial intelligence on employment, banking and transport sector. The following F ratios in ANOVA tests the independent variables statistically significantly influence the dependent variable. The above table revealed that the significance value ensures multiple regressions on the variables. The significance value is less than 0.05, therefore the impact of artificial intelligence which will influence to affect the various sectors like employment, banking and insurance sector. The multiple regression equation is on impact of artificial intelligence = 3.879 - 0.145(BS) - 0.166(TS) + 0.037(E). From the regression equation it is found that impact of artificial intelligence has more





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negative impact on transport sector. In similar way impact of artificial intelligence has next negative impact on Banking Sector. But impact of artificial intelligence has positive impact on employment. It may be concluded that the impact of artificial intelligence is more positively influenced to affect the future workforce and employment other than the banking and transport sector.

CONCLUSION AND RECOMMENDATION

Even though around 1.7 million jobs have been lost due to automation since 2000, according to the study, Artificial Intelligence is expected to create 97 million new jobs by 2025. Only a few people are aware about AI, so it can be made more accessible to every people in future. Possibly monotonous and the boring tasks will be given preference and replaced by the AI in future. Hazardous jobs that are being performed by human can be switched over to the AI, in this way unwanted loss of life can be prevented. By the implementation of AI in work force there will be less paper work, as it analyses the data and generates accurate and reliable data. Implementing AI- powered recruitment systems may lead to streamline the hiring process and reduce bias. By utilizing AI-based performance analysis tools to provide real-time feedback and insights, which can help employees improve their work quality. AI holds the key to unlocking a magnificent future where driven by data and computers that understand the world, and making more informed decisions. Artificial Intelligence (AI) has rapidly transformed from a futuristic concept to an everyday reality. As AI becomes more integrated into our daily lives, it has the potential to revolutionize industries and improve human experiences in numerous ways. Intervention of AI in the job sector has made an impact of doing the jobs effectively and with minimum error. Though the respondents have concern about their jobs being replaced and their privacy being disturbed by AI, they still choose AI for data related services as it cannot be altered or tampered to their needs. Artificial Intelligence will be unbiased and will ensure data accuracy.

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Table 1 Showing Chi-Square Test

Variable	Chi-Square p value	df	Sig. Value	S/NS	Remarks
Educational Qualification and Problems faced due to intervention of AI	0.111782	9	0.05	NS	Accepted

Table 2 Showing Multiple Correlation Co- efficient

Model	R	R Square
1	0.476	0.227

Table 3 Showing ANOVA test

F	Significance
1.667	0.042

Table 4 Showing Regression Co-efficients

Categories	Unstandardized Co-efficients (B)	Т	Significance
Constant	3.879	3.906	0.000
Banking Sector	-0.145	0.950	0.050
Transport Sector	-0.166	0.075	0.008
Employment	0.037	0.355	0.078





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RESEARCH ARTICLE

An Experimental Analysis for Forgery Detection of Handwritten Signatures using Deep Learning

Darshini Vipinchandran¹ and Jency Jose²

¹Lecturer, Department of Computer Science, Mount Carmel College, (Affiliated to Bangalore City University), Bengaluru, Karnataka, India.

²Assistant Professor, Department of Computer Science, Mount Carmel College, Bengaluru, Bengaluru, Karnataka, India.

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*Address for Correspondence Darshini Vipinchandran

Lecturer,

Department of Computer Science,

Mount Carmel College, (Affiliated to Bangalore City University),

Bengaluru, Karnataka, India.

Email: darshini.vipinchandran11@gmail.com



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ABSTRACT

Handwritten signatures are being used commonly for various transactions which include the banking sector, official documents, registration documents, and many more. It is very crucial to verify signatures as forged signatures can lead to the loss of personal and professional data. There are several ways of signature forgery which include random forgery, unskilled forgery, and skilled forgery. It is very difficult to distinguish between the original and the forged signatures when it is done by a skilled forgery expert. Hence the demand for signature authentication has increased as it is important for ensuring security and improving transparency. In this paper, a study on the different forgery detection methods of handwritten signatures is carried out, and is concluded that Convolutional Neural Network (CNN) is the most popular model used for datasets consisting of handwritten signatures. In this model, the signatures dataset is trained, and estimates are made to identify if the signature is authentic or forged. A survey on the various CNN architectures such as ResNet, GoogLeNet, AlexNet, and YOLO-v3 for forgery detection is carried out. The accuracies obtained using CNN is 96.69%, CNN with ResNet is 88.36%, GoogLeNet is 77.5%, CNN with AlexNet is 85%, and CNN with YOLO-v3 is 98.35%.

Keywords: Architecture, Artificial Intelligence, Convolutional Neural Network, Deep Learning, Detect, Forgery Detection, Handwritten, Machine Learning, Signature





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INTRODUCTION

Biometric systems are used to verify individuals as personnel identification is critical in organizations. Various biometric techniques are thumb print, iris examining, and signature verification, the latter of which is still acceptable for social and legal verification. Since everyone's handwriting is different, signature authentication is one of the processes of validating or detecting individuals[1]. An auto graph is defined as a distinctively carved illustration that an individual makes on a record. It is generally used to sign a cheque, a legitimate paper, or a contract. Any individual's signature portrays animage expressing a certain pattern of the pixel that belongs to a specific being. The problem arises when someone attempts to recreate it[2]. Since signatures are so widely used, many mischievous groups attempt to forge the signature for personal gain[3]. Images are used in almost every field, including health systems, news sites, social networking sites, legal investigations, schooling, armed forces, and firms. Swift technological improvements have led to the formation and distribution of a massive amount of images in recent years[4]. Forgery is a legal concept that refers to the objective to defraud someone. A forged signature has been proven to be false[3]. Image tampering has surfaced as a severe illegal issue[5]. Therefore, excellent signature forgery detection approaches are essential[3]. The key goal of signature detection is to sense the spot of signatures on challenging visualized documents and crop the required area. Present signature detection methods are categorized into two. One approach would be to recommend a procedure to extract characteristics and detect signatures. The further approach is to demonstrate the signature discovery task and perform it using deep learning-based methods[6]. Artificial Intelligence(AI) is a program capable of sensing, reasoning, acting, and adapting[7]. Machine learning(ML)is commonly considered a subfield of AI. It is the study of computer algorithms that assist systems to instinctively evolve and learn based on their experiences.

Machine learning algorithms enable machines to make sensible choices without the need for external aid. Such decisions are made by identifying significant underlying patterns in large amounts of complex data. There are three main types of machine learning algorithms depending on their learning technique, the kind of data they request and respond to, and the kind of difficulty they resolve. These are supervised, unsupervised, and reinforcement learning[8]. Deep learning(DL) is a subset of ML which in turn is a field under AI as seen in Figure 1. Deep learning is the procedure that uses multi-layer neural networks as seen in Figure 2 to integrate various machine learning algorithms. For understanding the input data, these distinct processing layers learn data representations with several levels of abstraction[8]. Furthermore, DL has progressively turned into the utmost extensively held computational methodology in the field of machine learning, accomplishing outstanding outcomes on a variety of complex cognitive tasks, identical to or as well outperforming human implementation[9]. Deep Learning learns from a massive amount of data. DL has grown rapidly and has many applications. Additionally, in domains such as cyber security, bioinformatics, automation and control, natural language processing, and health information processing, DL has outpaced well-known ML techniques[7]. The most well-identified deep learning networks are recursive neural networks (RvNNs), RNNs, and CNNs. Convolutional neural network (CNN) is the most well-known DL network. CNN's advantage is that without any human intervention, it can automatically identify significant features. In domains such as computer vision, speech handling, and face identification CNNs are widely used. CNNs were inspired by human and animal brain neurons[7].A CNN is made up of many convolution layers that come before sub-sampling (pooling) layers, and the final layers are Fully Connected layers as depicted in Figure 3. There are different CNN architectures namely AlexNet, ResNet, GoogLeNet, VGG, and DenseNet[7]. Conventional forgery techniques detect forgeries by aiming at several artifacts for instance changes in lighting, dissimilarity, compression, radar noise, and shadow present in a forged image. Each output map in CNN is created by sharing weights in a convolution operation. Furthermore, instead of relying on engineered characteristics to uncover precise forgeries, CNN uses learned features from training images and can reveal previously unnoticed forgeries. As a result of these gains, CNN is an excellent tool for detecting forgery in images. The CNN-based model can be trained to recognize the abundant objects present in a forged image[10].





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LITERATURE SURVEY

The review of the literature as seen in Table 1 provides a summary of existing research on machine-learning methods for detecting forgery in handwritten signatures. The review includes a description of the several algorithms, approaches, and CNN architectures used on different signature datasets. Jerome Gideon et al.[11]analyzed the image dataset of 6000 images. As CNN is the best algorithm for the image dataset, the authors have considered the CNN algorithm in their paper. They got three different accuracies by splitting data into three different training and testing ratios which are 8:2, 7:3, and 6:4, where the 8:2 split gave a better performance with 98.23% accuracy. The proposed system was successful in identifying if the signature was forged or not. However, the structure of the fully connected layer is not optimal[11]. Khan et al.[12] took two datasets into consideration, one with 10,868 records and another one with 3000 benign files. By using two different architectures of CNN, namely GoogLeNet and ResNet the authors were able to construct a model. ResNet model outperformed GoogleNet by showing high accuracy of 88.36%[12]. Jahandad et al.[1] have utilized GoogLeNet's Inception-v1 and Inception-v3 algorithms on the GPDS signature dataset. After analyzing and comparing both models the authors noticed that Inception-v1 showed a better performance with an accuracy of 83%. By considering data augmentation as a step for pre-processing, the overall performance of the model and over fitting of data can also be avoided in future work[1]. Poddar et al.[2]proposed a system that was able to successfully identify the signature holder by checking if the signature is forged or not. The system was economical in recognizing forgeries at the runtime. The authors considered a dataset with 1320 images and made use of the CNN, Crest-Trough method, and SURF algorithm & Harris corner detection algorithms. Among which Harris Algorithm gave the highest accuracy of 94%. In future work, the resulting approach accuracy can be increased by attempting different and better parameter coefficients that increase the deviation between real and forged signatures[2]. Kao et al.[13] have collected data from ICDAR-2011 SigComp for forgery detection.

As the author is detecting forgery using the image dataset the author has used the CNN model which is one of the best-performed algorithms for the image dataset. The algorithm has performed better than average, with an accuracy range of 94.37% to 99.96%. The findings determine that the network's performance may be considerably enhanced by raising the numeral of falsified sample seven with a single known sample[13]. Abdulhussien et al.[14] analyzed the dataset which was collected from SID Arabic handwritten signatures, CEDAR, and UTSIG. Using the One-Class SVM(OCSVM) algorithm, the model was built for the collected dataset. "The system improved the 10% false acceptance rate (FAR) of Arabic signature verification on the SID-Arabic signature database without affecting computation time, which achieved 0.037 false rejection rate (FRR), 0.039 FAR skilled, 0.063 FAR simple, and 0.044 and equal error rate (EER). For the UTSig and CEDAR, the suggested method produced EER values of 0.074 and 0.048, respectively". This experiment presented a higher peak signal-to-noise ratio(PSNR) and a lower mean-square error(MSE)[14]. Lopes et al.[15] to verify handwritten signatures, employed a dataset from CIFAR-10. To confirm the handwritten signatures, AlexNet Architecture is been used. The author stated that the model provides a highaccuracy classification for a limited number of samples. The author discussed the study's shortcomings and noted that it takes a long time for AlexNet to train a new model. The described model might perform better if local feature extraction is added[15]. Ryan et al.[16] utilized the YOLO-v3 architecture of CNN on 300 images which were split into an 8:2 ratio. This architecture improved the performance and ability of signature forged detection. This system gave an accuracy of 98.35%. However, new feature selection methods can be implemented to improve the effectiveness of the model[16].

CONCLUSION

Thus the paper focuses on the forgery detection of signatures. Dependable and accurate signature recognition and verification systems are critical for many applications, including law enforcement, access controls, and many corporate processes. It can be used as a middleman to validate multiple reports such as cheques, official records, and certificates. CNN and its different architectures are implemented in most of the papers as it is the best model for





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datasets with images in which the signatures datasets are trained, and estimates are made to identify if the signature is authentic or forged. The model produced promising results when CNN architectures were considered. In the future, the accuracy of forgery detection on signatures can be enhanced by increasing the layers in the CNN architectures or by introducing ensemble hybrid models. Due to the difficulty in identifying skilled forgeries in signatures, authentication based on biometrics is significant along with handwritten signatures.

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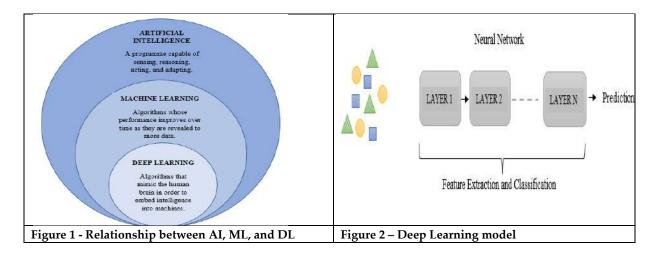




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Table 1 – Comparative Analysis

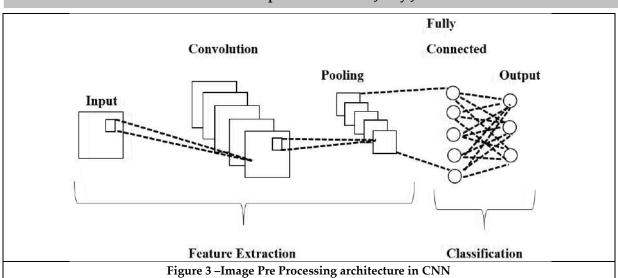
Author Name	Dataset	Algorithm	Accuracy/Result
S Jerom Gideon et al.	6000 images	CNN with 8:2 split	98.23%
		CNN with 7:3 split	94.83%
		CNN with 6:4 split	96.75%
Riaz Ullah Khan et al.	1) 10,868 records	GoogLeNet	74.5%
	2) 3000 benign files		
		RestNet	88.36%
Jahandad et al.	GPDS signature	GoogLeNet-	83%
		Inception-v1	
		GoogLeNet-	75%
		Inception-v2	
Jivesh Poddar et al.	1320 images	Harris Algorithm	94%
		Surf Algorithm	85%-89%
Hsin-Hsiung Kao et al.	ICDAR-2011 SigComp	CNN	94%-99.96%
Ansam A et al.	SID Arabic handwritten	SVM	The experiment
	signatures, CEDAR, and		provided high peak
	UTSIG		signal-to-noise ratio
			(PSNR) and a low
			mean-square error
			(MSE)
Jose A et al.	CIFAR-10	AlexNet	85%
Ryan C et al.	300 photos	(You Only Look	98.35%
		Once) YOLOv3 of	
		CNN	







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RESEARCH ARTICLE

Real-Time Facial Recognition using Drone

Arunachalam. R1* and Poovarasan Selvaraj2

¹UG Student, Department of Information Technology, Sri Ramakrishna College of Arts & Science, Coimbatore, Tamil Nadu, India.

²Assistant professor, Department of Information Technology, Sri Ramakrishna College of Arts & Science, Coimbatore, Tamil Nadu, India.

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*Address for Correspondence Arunachalam. R

UG Student,

Department of Information Technology,

Sri Ramakrishna College of Arts & Science,

Coimbatore, Tamil Nadu, India.

Email: 21107071@srcas.ac.in



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ABSTRACT

The study develops a facial recognition Unmanned Aerial Vehicle (UAV) solution to help the task force identify criminals, missing people, civilians, and for surveillance purposes. One type of technology that requires knowledge of face detection and identification is a facial recognition system. Typically, it's employed to authenticate users by extracting face traits from photos and confirming user identities. This paper describes an unmanned aerial vehicle (UAV) equipped with a camera, linked to facial recognition software, and utilized for remote robot control. With an accuracy of 98.6% when the drone's camera angle is within 37 degrees, the proposed approach can adapt to changing drone locations. Certain applications help law enforcement officers with their investigations, such as crowd monitoring, crime analysis, search and rescue, and incident investigations.

Keywords: Drone, Facial Recognition, LBPH, Face Net

INTRODUCTION

Technology for facial recognition has gone a long way and is currently developing quickly. We need to understand convolution neural networks (CNN) in order to comprehend the facial recognition algorithms employed in this thesis. Since the 1980s, CNNs have existed and are nothing new. They belong to a class of deep learning methods that may be used to give weights, biases, and learnable significance to different characteristics or objects in a picture. Then, one item may be distinguished from another using these assigned traits. Using convolutional layers, a CNN can recognize an item in a picture. Filters in these layers are capable of detecting simple patterns like an edge or a line. More accurate filters may be applied when convolutional layers are added. A CNN may move from identifying





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edges and lines to forms and then to objects, persons, or whatever else it is intended for by iteratively cycling through the filtering process. This technique may now be used considerably more swiftly and effectively, for example in conjunction with a live video broadcast, since processing power keeps increasing. The primary technique for facial recognition systems is now CNNs. Algorithms for facial identification and detection can make use of prebuilt models that have been trained using thousands of photos and countless hours of computation. In this thesis, pre-trained models will be combined with the chosen face detection and identification algorithms to develop a drone that can recognize and identify people as well as follow their movements. While computing power has increased, the majority of CNNs still require high-end graphics processing units (GPUs), which are too expensive and heavy to be used in lightweight, low-cost drones. The drone's photos can be sent to a computer with enough processing capacity to help solve this problem. Drones use 2.4 gigahertz radio waves, which are widely compatible with Wi-Fi, to communicate with external devices. This network may be used by the computer attached to the drone to exchange information with it. Faster processing speeds and far better drone reaction times for tracking will result from using a considerably more potent technology for picture analysis. The network does have limitations, though. There is a potential that the drone and computer might lose contact for brief periods of time or perhaps become completely separated. The computer and drone have developed a network that allows for two-way communication.

Real-time instructions may be sent from the computer to the drone to control its movement while the drone can provide a live video feed to the computer for analysis. The drone may also update the computer with status information, such as battery life, flying time, and current altitude. Without input from the computer, the drone won't know what to do, hence this network is essential for the system to work. The greatest network range of a drone is a crucial factor since disconnected systems tend to malfunction. The facial tracking component is the last element of the system used in this thesis. The software allows for the designation of a single person as the target, and when that person is located with a confidence level of at least 80%, a specific flag is set alerting the tracking component of the system. The region of the target person's face in the frame in which they were recognized is calculated during tracking. A tiny region denotes a distance from the objective, whereas a big area denotes proximity. In order to keep the target inside a given range in relation to the drone, the computer will perform this analysis and transmit movement directions to the drone. Because the target may swivel their face away from the drone, the drone can also identify the person's centre of the face. The optimal movement instructions will be found and relayed by the computer if the target turns away from the drone; research has shown that this is the best posture for facial identification and recognition. It is crucial for the system to swiftly examine frames since the tracking algorithm only transmits instructions for one frame at a time. One or more of the facial detection or identification systems may operate slowly, review frames from previous seconds, and transmit inaccurate commands to the drone if this happens.

RELATED WORK

Drone has been somewhat of a buzzword recently as interest in them continues to increase and associated prices appear to be falling. Many people have been able to conduct research and carry out computer vision-related tasks while using drones as a result of this accessibility. The major ideas of this thesis are around using a drone to create a system that does real-time facial tracking and recognition. Despite the fact that other academics have carried out related work, none of them have put forth the same real-time automatic face recognition system that tracks people using drones as this thesis does. Academia Sinica's Institute of Information Science researchers have developed a drone-based facial recognition system. The effectiveness of drone face identification was tested empirically using two facial recognition systems—Face++ and identification—while taking into account variables including distance from a person, angle of depression, and drone height. Height, distance, and angle of depression all correspond to the location of the drone. As the drone ascended to heights of 3, 4, and 5 meters, pictures were captured. The drone progressively backed up to a distance of 17 meters from the subject while taking pictures every.5 meters starting at a distance of 2 meters. The researchers discovered that accuracy significantly declined as drone height and proximity to the person grew. At 1.5m for the drone height and 2m for the drone distance from the person, Face++ and Recognition both had 100% accuracy. However, the accuracy of both facial recognition algorithms substantially





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decreased as subject distance and drone height grew.17-meter distance Averaged by Face++[5], 5% accuracy at drone heights between 2 and 5 meters. For the same distance, Recognition's accuracy averaged 5%, a significant gain but still far from adequate. Other noteworthy difficulties were scenarios in which the drone was high in the air yet there was little space between it and the subject. Such circumstances led to a high angle of depression, which decreased accuracy and increased the number of true positives—where a person was mistakenly recognized as someone else. This study demonstrated the viability of using drones and face recognition systems together, but the findings also highlighted the system's many drawbacks, which might compromise its accuracy. Researchers from the National Research Council's Institute of Information Science and Technologies (ISTI- CNR) in Pisa also set out to investigate the barriers preventing the use of facial recognition in combination with drones. The key limitation they took into account was the possibility of humans in the picture appearing in low resolution with a drone at a high height.

As a low-quality image makes it difficult to distinguish objects in the frame, this would impair facial detection and recognition systems, thus they sought for ways to enhance it. For facial recognition, they used three models: ResNet-50, SENet, and their own model. They employed dlib, MTCNN, and Open CV-DNN for facial detection. The Drone dataset, which includes both low- and high-quality photos, was used to test and evaluate all three algorithms. During testing, a low-resolution image was used, a facial detection system was applied to it, and then a facial recognition system was used to compare it to its high-resolution counterpart. The top performing system, SENet, only had an accuracy of 24.25 percent utilizing MTCNN at the beginning. They reran the experiments with tighter cropped faces that excluded more of the background, speculating that the poor accuracy was due to the cropping of the detected faces. The model they created with an accuracy of 60.87% utilizing MTCNN was the best-performing system, and the results significantly improved. The efficiency of SENet has also significantly increased; it currently reports an accuracy of 53.56%[1]. This study demonstrated the significance of the identified face's cropping in lowquality photos and the possibility that a tighter crop might improve accuracy. Other researchers at the Xi'an Aeronautical University developed a drone-based security system employing an LBPH face recognizer. The facial recognition technology was operated by the researchers using a Raspberry Pi mounted to their drone. A trained LBPH facial recognizer was employed to clip those faces after the system utilized Haar Cascade to find them in the crowd. The user would be notified if the facial recognizer discovered a match in its dataset so they could take over physical control of the drone or give it fresh instructions. 250 photos of each individual were used to train the facial recognizer, and each image was cropped to solely contain the person's face. To reduce the quantity of information in the photo and shorten the training period for the model, the cropped image was downsized and made black and white. The algorithm could positively identify a person with a minimal degree of confidence of 50%. This minimal degree of confidence was chosen since several problems, particularly those relating to distance, surfaced during testing. The recognition system often found people but lacked the confidence to classify them. To enhance recognition while minimizing false positives, the confidence threshold was gradually lowered.

With an overall accuracy of 89.1%[11], this system was able to recognize individuals in a crowd. For their facial recognition system, researchers at The LNM Institute of Information Technology also went to LBPH. They used LBPH and FaceNet, two facial recognition algorithms. They put these technologies into use on a Raspberry Pi and connected it to a drone, much like many other researchers. It is not a very powerful gadget, but having a Raspberry Pi on a drone has several benefits because all essential processing can be done on it. The drone was able to pinpoint a specific target in a crowd on its own using the two face recognition algorithms. The system did work despite a number of problems, but they determined that because of the hardware restrictions of the Raspberry Pi[7], building an accurate model that could distinguish a face from a crowd is a very difficult undertaking. By transmitting the drone's video to a computer with the appropriate processing ability to precisely analyse each frame and transmit data back to the drone, the technique used in this thesis will address that problem. An extremely comparable experiment was carried out by researchers at the RMK College of Engineering and Technology in Chennai. Additionally, they employed LBPH for facial recognition and Haar Cascade to detect faces in their system. They claimed a far higher accuracy percentage of 98.6%[6], but their system would check the collected faces against every single test image accessible, which could account for this high accuracy rate. The disadvantage of this strategy was that it caused the system to operate at a very sluggish processing speed of just 9 FPS. Both the researchers at Xi'an





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Aeronautical University and the researchers at the RMK College of Engineering and Technology employed the identical LBPH system. In addition, LBPH was used in this thesis along with Haar Cascade for the system's facial recognition component. The difference between this thesis and the method developed by the researchers at Xi'an Aeronautical University is that when a certain face is found, the drone will independently start tracking the person instead of giving that information back to the user and waiting for feedback. Deep Drone is a technology that Stanford University researchers created for tracking and identifying objects. In contrast to the method in this thesis, the authors' paper primarily focused on identifying and tracking general populations rather than a single individual. Two basic systems make up Deep Drone; the first is a detection technique that uses a CNN to look for any individuals in the frame. The second system is the tracking algorithm, which makes use of kernel correlation filter (KCF) and histogram of oriented gradient (HOG) characteristics. The device could only scan at 1.6 FPS due to the highly slow and computationally demanding process of detecting a human.

When a person was found, the system would draw a box around them and provide the tracking algorithm the location. KCF was able to process much faster at 71 frames per second since it was only conducting tracking on the frames [4]. This method has the disadvantage that it must switch back to utilizing the detection algorithm in order to find people if the subject moves too quickly or if another event causes the tracking algorithm to lose sight of the target. They overcame problems with side profiles and targets that were continually moving by decreasing the amount of confidence needed to designate an object as a person. Although lowering the confidence level might seem like a simple remedy, it can lead to more false positive matches. Modern technologies on the drone were also operating too slowly, making it impossible for it to send movement commands rapidly enough to track the target. They ultimately chose to employ the KCF tracking system because it needed less processing power, but this older method did reduce accuracy. Deep Drone is designed to find and track any person that it finds, however the technology used in this thesis is not. The system used in this study will, however, have a single target person identified in the program, and it will independently discover and track only one person, addressed by scholars at North Carolina State University, Raleigh. They primarily addressed problems discovered by researchers at the Institute of Information Science, Academia Sinica, who discovered that the height of the drone was a significant role in facial recognition. The researchers employed three models-VGG16, VGG19, and InceptionResNetV2-for their investigation. These models were chosen because they are among of the most precise ones in use right now.

They used the Drone Face dataset, which was created to help improve current drone capabilities, to train these algorithms. These pictures were taken from various heights, ranging from 1.5 to 5 meters, and at various separations, ranging from 2 to 17 meters from the subject. The 3 models were then trained using these photos; however, each model underwent a distinct training process for each test. The goal of the testing was to identify the facial recognition model that could adapt to changes in drone altitude the best. The models were tested on photographs taken at 5 meters of drone height after being trained on images acquired between 0 and 4 meters of drone height. When the drone's height was higher during training than during tests, the researchers discovered that models were more accurate. They did discover, however, that InceptionResNetV2 and VGG16 had the ability to recognize faces at a height greater than the height they were trained on. Additionally, they discovered that when the height varied, identification activities were far less influenced than verification ones. When trained on heights lower than the height of the test images, all 3 systems-maintained accuracy levels of over 90% during recognition tasks, but only between 34 and 38% during validation tests[2]. They found that training the model with images that are taken from a greater distance and at a higher drone height helped to increase the accuracy of their systems. Additionally, they discovered that InceptionResNetV2 provided the most reliable and accurate findings.

METHODOLOGY

The methodology portion will cover the implementation frameworks for the system created in this thesis. The hardware will be operated by a drone, which will send video from its camera to a computer. This computer will look through each frame of the video feed in real time for a specific person using facial recognition and detection





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algorithms. If the person is found, the system's tracking function kicks in, and the drone is given movement instructions by the computer to keep the intended subject in the picture. When these little systems work together, they can form an autonomous drone that is able to find and follow a specific individual.

Drone Operation

In recent years, as drones have become more readily accessible, their use has increased. Today, individuals use drone technology in previously unthinkable ways, such as package delivery, tree planting, and even swiftly providing medical supplies to those in need. The frame, motors, propellers, electric motor controller (ESC), power distribution board (PDB), flight controller, battery, receiver, camera, video transmitter (VTX), and sensors are the 11 fundamental parts that make up a drone. A drone has four degrees of freedom as well. Three translations and one rotation make up the four degrees. Drones can move up, down, left, right, and forward and backward thanks to the three translations. Drones can rotate both clockwise and counter clockwise using this single revolution. Height, distance, and angle of depression are taken into consideration when using the drone. A tiny drone was employed for this thesis. It is a light drone that only weighs 80 grams but has a 13-minute flight time, a 100-meter range, and a 5-megapixel camera that can provide HD video in 720P. This project also made use of the djitellopy Python library, which made it simple to transmit instructions to the drone and receive data from it directly to a computer. A link might be made between the drone and a computer using the connect() function. The stream on() function then transfers the video recorded by the drone to a computer so that it may be used to perform facial recognition on the image. The facial recognition algorithm examines the footage pixel by frame even though it is being streamed live.

The system can also show the stream on a computer and give more details if any individuals are recognized. The tracking component of this project was implemented using the djitellopy library's numerous distinct functions for regulating drone movement. When a target is located, the system allows the user to identify a person as a target, and the drone will then follow that person. A box is created around each face to show who the algorithm has identified it as photographs are evaluated and persons are identified. The system issues movement commands based on the size of the box drawn around the target, if one of those recognized individuals is the target. The drone will get closer until it reaches a predetermined range if the box's surface area is shrinking or has already shrunk. If the area is too large or the target is approaching, the same principle applies. If the target moves side to side, similar movement commands are utilized to rotate the drone, ensuring that the target can be tracked. However, for best tracking, the target's frontal face should always be visible. As indicated in the results section, side profile photographs result in decreased accuracy and images of the back of the head result in zero accuracy.

Facial Recognition Systems

The majority of facial recognition technologies now operate as depicted in Figure 3.4. The system starts with an input image and looks for any faces. If any faces are discovered, they are cropped, and information about facial features is extracted to produce a face embedding, a vector that can be used to identify a person. A classifier is then given access to that embedding or the face traits. The classifier is a model that has already been pre-trained and contains data on one or more individuals that it was taught to categorize. The classifier can output details about the person(s) recognized using the input data and any minimum confidence threshold value set. The three systems that were used in this thesis all function largely similarly. To verify the functionality of the system used in this thesis, three distinct facial detection and recognition systems were constructed for this research. The systems of preference were LBPH face recognizer, FaceNet, and Face Recognition. The LBPH face recognizer was selected because it has been used in earlier work with drones and because it provides a baseline for findings. Due to their remarkable performance in the Labelled Faces in the Wild (LFW) benchmark dataset, FaceNet and Face Recognition were chosen.LFW is a widely accessible dataset that is used to assess a system's face verification accuracy. FaceNet and face recognition both reported 99.38% and 99.63% accuracy, respectively, using the dataset. A test dataset was used by all three systems to train their classifiers to recognize certain individuals. Each participant has 15 tagged photos in the test dataset. Five pictures were shot in a low light environment, five pictures in an ambient light setting, and five pictures in a high light setting. All of the photos were taken with a frontal or side perspective of a person's face against a simple background. On the photos that were given to each system, some uniform actions were also carried





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out. All of the offered test photographs were scaled and made black and white for training purposes. Additionally, each photo's face was identified by the algorithms, which then cropped it so that only that portion of the image was used for training. However, because each approach gives certain parts of the face a distinct relevance value, none of the three algorithms produced the same cropping of a face. Streaming photographs from the drone to the computer is another consistent operation. Each system resizes the image and converts it to black and white in order to speed up the facial recognition process because the photos transmitted by the drone are very huge and bright. Each system also shows the stream to the user in a window. To alert the user that a facial detection has taken place, every system draws a box around each face it has identified in the frame. A name will be added to the box if a face matches one in the learned classifier; otherwise, it will simply say "unknown." The program's tracking feature will turn on whenever a certain target is located.

Implementation of LBPH

Due to the volume of prior studies that have used this technology, LBPH face recognizer was chosen. When using LBPH, a pixel is chosen along with a collection of pixels in a predetermined radius. A histogram is then produced using the data collected, and a regional level histogram is produced by combining the pixel level histograms. In order to create a histogram that provides a general description of the identified face[8], the regional level histograms are finally concatenated. To locate similar histograms that would aid in identifying a person, this general histogram can then be utilized for comparison with a classifier. The four basic LBPH parameters that are frequently used are radius, neighbors, grid X, and grid Y. The radius, which indicates the area surrounding the central pixel in the circular local binary pattern, is often set to 1. The number of sample points called "neighbors" is often set to 8 when creating a circular local binary pattern. The numbers of cells in the horizontal and vertical planes are represented, respectively, by grids X and Y. This value is typically set to 8, and more cells produce a finer grid with a larger dimensionality for the final feature vector. Making the model is the first stage in putting the LBPH system into practice. Haar Cascade, a facial detection technique first introduced in 2001 [10], is employed to help create the model. A cascade of classifiers with ever higher levels of complexity is how this system operates. The frontal face cascade, which has classifiers that are excellent at frontal face detection, was employed in this study. LBPH face recognizer and Haar Cascade are combined to construct the trained model. The provided annotated photos are first converted to grayscale by this system before Haar Cascade is used to find faces in each image. After being clipped from the image, the discovered faces are then passed to the LBPH face recognizer, which uses them as training images to build a model. The facial recognition system is then updated with the model. In this setup, a drone continuously broadcasts live video that is dissected and evaluated frame by frame. The trained LBPH face recognizer algorithm may then be used to run and return any matches after using the Haar Cascade to detect any faces in each frame.

Implementation of FaceNet

A universal embedding system called FaceNet has been put out by Google researchers [9]. It functions by employing a deep convolutional network to embed a picture in a Euclidean space. These embeddings include 128 face measures in them, which are subsequently translated to a Euclidean space where like images are grouped together. Although the network's methodology for constructing the embedding is unknown, the results stand on their own. The technique locates the Euclidean embedding of an unknown image and categorizes it according to the nearest known individual. FaceNet was implemented using Tensor Flow for the sake of this thesis. There are three main phases to implementing FaceNet. The first step is pre-processing, which involves cropping and aligning each face in the training data set. The following phase entails learning or embedding face representations in multi-dimensional spaces where distance correlates to the indicator of face similarity. An embedding and classifier were made using the training dataset. The classifier, which integrates the FaceNet model with given Tensor Flow, detect face function, is the last step. Each frame that the drone streams to the program is shrunk, turned to black and white, and then inspected. A Multitask Cascaded Convolutional Neural Network (MTCNN) is built using the Detect face function to locate faces in a frame and store them in an array. The three stages of MTCNN's operation are the proposal network, refine network, and output network. Before the first stage, a preliminary process is carried out. The image is first





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scaled up and down several times to help find faces of different sizes. After the preliminary procedure is finished, the proposal network stage is launched. The proposal network is made up of numerous smaller processes, and its first process uses a very low threshold to accomplish facial detection. Due to the low threshold for facial detection, any object in the frame that even somewhat resembles a face will be classified as one. Although this technique yields a lot of false positives, it ensures that all potential faces in the image are recognized. The generated regions constitute the "proposal" and are forwarded to the refine network stage, which is the next step. This stage involves filtering the area and building larger, more valuable bound boxes around the detected faces. The output of the refining network is then used as the input for the output network, which is the last stage. The bounding boxes are final refined in this step, producing a very accurate output of the detected face. The person's identification is then returned once the photos are mapped to the trained classifier and the minimum threshold value is reached

Implementation of Face Recognition

The "world's simplest face recognition library" is called face recognition[3]. Face Recognition is constructed using the deep learning face recognition algorithms from dlib, a toolbox for building complicated software with machine learning techniques. In order to find faces in an image, Face Recognition employs the histogram of oriented gradient (HOG). HOG works by choosing a pixel in the image and examining its neighbours to determine how dark the chosen pixel is relative to its neighbors and which direction the image gets darker.[11] After repeating the process for every single pixel, the program keeps track of which way the image becomes darker, and the result is a gradient that depicts the movement of light throughout the entire picture. The direction of each square is chosen depending on the direction that is most common in that part after the image has been divided into 16 by 16 squares. The HOG face pattern, developed by training on numerous faces, is then applied to this version of the image to determine whether or not a face is present in the image and where it is located. face recognition centres and aligns its photos, just like FaceNet does. It maps 68 precise spots on the face using a facial landmark estimation algorithm, which enables the program to centre or realign the image as necessary, face recognition extracts a 128-measurement embedding of each face from the training data set using a deep CNN. The Google researchers that developed FaceNet also developed this system. Encodings were created once the test data set was loaded for the face recognition implementation. Figure 3.6 provides a summary of the actions the face recognition system takes to identify and detect faces. It reads the incoming video stream frame by frame, resizes the frames, and converts them to black and white, just like the other systems. Face encoding was then utilized to create an embedding for each face that was discovered after using HOG to find faces in the frame. These embeddings were then put in comparison with the known face embedding, and the outcome was given. Compared to LBPH or FaceNet, face recognition was implemented much more quickly and efficiently. This is due to the Python face recognition package, which includes functions for many of the implementation processes outlined above. Most of the functions in LBPH and FaceNet have to be written from scratch.

CONCLUSION

The paper describes the development and implementation of a real-time facial recognition and tracking system that uses a drone and computer to find and track a predefined target individual. The facial recognition systems were found to be the most important component of the entire system, and their performance was measured using three separate techniques. Even though each of the other facial recognition systems had unique advantages and disadvantages, FaceNet fared the best when tested in an actual scenario. The results of this study suggest that using a drone in conjunction with face recognition software makes it possible to track an individual. By using a lightweight, inexpensive drone to send footage to a computer that can recognize faces, facial identification is still possible. By streaming video to a computer, this method eliminated a lot of the performance issues found in papers that constructed related systems. Although this system's capabilities are limited for unconnected reasons, it provided a strong foundation for further development.

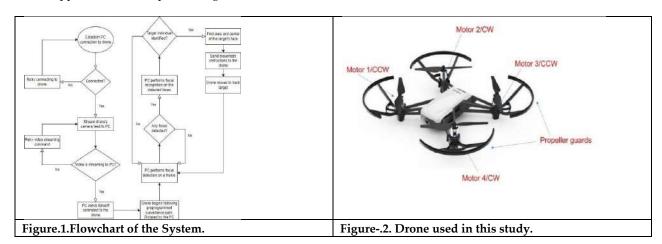




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REVIEW ARTICLE

A Review on Enhancing Educational Outcomes: Leveraging Educational **Data Mining for Predictive Analysis**

D. Gajalakshmi^{1*}, P.V.Praveen Sundar² and R. Denis³

¹Research Scholar, Adhiparasakthi College of Arts and Science, Kalavai, Ranipet Dt,(Affiliated to Thiruvalluvar University, Vellore) Tamil Nadu, India.

²Assistant Professor, Adhiparasakthi College of Arts and Science, Kalavai, Ranipet Dt,(Affiliated to Thiruvalluvar University, Vellore) Tamil Nadu, India.

³Department of Computer Science, Mount Carmel College (Autonomous), Vasanth Nagar, (Affiliated to Bangalore City University), Bengaluru, Karnataka, India

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*Address for Correspondence

D. Gajalakshmi

Research Scholar, Adhiparasakthi College of Arts and Science, Kalavai, Ranipet Dt, (Affiliated to Thiruvalluvar University, Vellore) Tamil Nadu, India.



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ABSTRACT

Data mining techniques have gained significant interest in the field of education, particularly in educational data mining (EDM). EDM is the application of conventional data mining techniques to address issues related to education, such as student demographics, academic transcripts, test scores, attendance, and student queries. EDM has become a potent tool for predicting academic accomplishment, finding hidden patterns in educational data, and enhancing the learning and teaching environment. The subfields of Data science and machine learning have proven to be extremely wellorganized and significant in many domains, including education, over time. A computer system can collect data, evaluate it, and make inferences that appreciate to a feature of artificial intelligence known as machine learning. Assessment systems that predict student achievement have recently been developed in the education sector using data mining and machine learning techniques. Machine learning techniques are being researched in the realm of education to identify meaningful patterns that improve student comprehension. Based on those patterns, academic institutions are making decisions. One important educational measurement metric that affects institution accreditation is the assessment of student performance. Underachievers in those colleges need to be counseled in order to implement a plan for raising student performance. It helps teachers and students work through problems that come up because of the students' coursework and the teachers' teaching strategies. In terms of predicting student performance, this review study looks at a number of studies related to detecting low-performing pupils.





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The results of the survey indicated that several machine learning techniques are used to address problems related to risk assessment and student performance evaluation. Machine learning approaches play a major role in improving the student performance prediction system and advancing the field of student performance prediction.

Keywords: Machine Learning(ML), Logistic regression, KNN, SVM, XGBoost, Naïve bayes

INTRODUCTION

Learning analytics has been given a new dimension by EDM, which encompasses various aspects of gathering student data, understanding the learning environment through examination and analysis, and revealing the best student/teacher performance. This includes the collection, measurement, and reporting of data about students and their contexts to optimize learning and its environments. EDM can predict student academic performance, identify trends in system access and navigational behavior, and identify and remove potentially failing students. Digital data left by learning management systems (LMS), student information systems (SIS), intelligent teaching systems (ITS), massive open online courses (MOOCs), and other web-based education systems can be analyzed to assess potential student behavior. These data can be used through EDM to analyze the activities of successful and at-risk students, create remediation plans based on students' academic performance, and support teachers in creating pedagogical approaches. EDM helps instructors forecast events such as dropping out of school or losing interest in the course, analyze internal factors affecting their performance, and develop statistical tools to predict students' academic achievement. Early prediction is a novel phenomenon that involves evaluation approaches to assist pupils by offering appropriate remedial measures and policies in this field. Learning management systems have become an integral aspect of higher education, particularly during the pandemic period. Universities should now strengthen their ability to use these data to forecast academic achievement and ensure student advancement. EDM provides fresh knowledge to educators by uncovering hidden patterns in educational data, allowing some areas of the education system to be assessed and improved to ensure educational quality.

Related Work

E-learning systems have been effectively analyzed in a number of EDM research (Lara et al., 2014). Some research have also attempted to classify educational data (Chakraborty et al., 2016), while others have attempted to predict student achievement (Fernandes et al., 2019) [1]. Asif et al. (2017) and Cruz-Jesus et al. (2020) used machine learning approaches to predict undergraduate students' academic performance. Asif et al. (2017) categorized students into low and excellent achievement groups, suggesting instructors should focus on a few courses with high or low performance to provide early warnings, support, and opportunities [2]. Cruz-Jesus et al. (2020) used 16 attributes to predict academic achievement, using machine learning algorithms like random forest, logistic regression, k-nearest neighbors, and support vector machines [3]. Fernandes et al. (2019) developed a model using demographic information and academic achievement grades to predict students' academic progress. The model used the Gradient Boosting Machine (GBM) to estimate achievement scores and absenteeism. Other studies identified students at risk of failing based on registration data and environmental factors. Hof-faith and Schyns (2017) used DM methods to diagnose students with probable issues and rank them based on their level of danger [4]. Rebai et al. (2020) used machine learning to identify factors influencing school academic performance, finding school size, competition, class size, parental pressure, and gender proportions as the most relevant characteristics. The random forest algorithm revealed that school size and percentage of girls significantly impacted the model's predicted accuracy [5]. Ahmad and Shahzadi (2018) and Musso et al. (2020) used machine learning to predict academic performance and dropout rates. Ahmad and Shahzadi's method, which uses learning ability, study habits, and academic interaction traits, had an 85% accuracy rate [6]. Musso et al.'s model, based on learning techniques, social support perception, motivation, socio-demographics, health status, and academic performance factors, predicted academic achievement and





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dropouts [7]. Waheed et al. (2020) developed a deep learning model based on student records for LMS navigation, showing that demographics and clickstream behaviors significantly impact student performance. They concluded that this model could be useful for early performance prediction [8]. Xu et al. (2019) found a correlation between internet usage habits and academic achievement, with internet traffic volume and frequency features positively correlated [9]. Bernacki et al. (2020) found that 75% of students needing to repeat a course were successfully predicted using a behavior-based prediction model. This strategy can help identify and assist students who might not succeed in the upcoming semesters [10]. Burgos et al. (2018) designed a tool for predicting future achievement grades and designed a tool for those likely to fail [11]. A survey of earlier studies looking to forecast academic success shows that a variety of machine learning techniques, such as neural networks, multiple, probit, and logistic regression, as well as C4.5 and J48 decision trees, have been used. Nonetheless, recent studies have used Naive Bayes algorithms (Ornelas & Ordonez, 2017) [12], genetic programming (Xing et al., 2015) [13], and random forests (Zabriskie et al., 2019) [14]. These models achieve extremely high prediction accuracy.

A thorough study of the elements and characteristics that affect students' performance and accomplishments is necessary for accurate prediction of academic performance (Alshanqiti & Namoun, 2020) [15]. In order to do this, Hellas et al. (2018) examined 357 publications on student performance that included information on the effects of 29 attributes. These characteristics were mostly associated with psychomotor skills, including performance in the course and prior to it, student participation, student demographics including gender, high school achievement, and selfcontrol. However, the primary factors influencing dropout rates were habits, lack of development, social and financial concerns, and career transitions among students [16]. According to Praveen et al. [17], identifying disengaged learners more precisely can be achieved using region-wise classification of online learners. The literature evaluation comes to the conclusion that a wide range of characteristics influence the ability to forecast a student's academic performance. Classifying pupils according to where they live also aids in the identification of disengaged learners. According to studies [17], [18], learning is insufficient on its own to identify disengagement. For improved prediction outcomes, they combine the time spent characteristic with the marks secured attribute. As was said in the literature evaluation, a great deal of important characteristics are utilized to forecast the student's log data. The academic performance of students was predicted by Praveen [19] using the naive Bayes method, and the Bayes model assisted teachers in identifying individuals who needed extra support. According to a survey on regional benchmarks of rural and urban students conducted by Praveen et al. [20], there is no significant difference between students who completed their previous coursework in a uniform area (rural or urban), but there is a significant difference between students who completed their coursework in a mixed area.

RESULT AND DISCUSSIONS

A Case Study Data Used

Here, the datasets taken from kaggle for the comparison of the best performing prediction algorithm among Logistic regression, KNN and support vector machine and other classification algorithms. There were a total of 791 rows of secondary students information with 31 columns of features. The data attribute include demographic social and school related features. Using the ratio of 0:30 split amongst training and testing instances.

Algorithm Used

Support Vector machine

A machine learning approach called SVM classification finds a hyper plane to divide several classes in a dataset. Support vectors, which are the closest data points to the hyper plane, are used to find the solution, which optimizes the margin between the classes. SVM is a memory-efficient, reliable algorithm that can handle datasets that are linearly and nonlinearly separable. Finding the hyperplane that divides the classes and optimizes the margin between them is the goal of the SVM classification algorithm. A two-dimensional linear SVM hyper plane has the following formula:







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Where the input vector is x, the bias term is b, and the weight vector is w. The decision boundary is given by the formula wT x + b = 0. The input vector belongs to one class if the equation's answer is positive; if it's negative, it belongs to the other class. Non-linear SVM classification is employed when the data in the original feature space cannot be split by a linear hyperplane. To move the data into a higher dimensional feature space where the classes might be linearly separable in this case, the SVM technique uses a kernel function. The Radial Basis Function (RBF) sigmoid, and polynomial are the three most often utilized kernel functions. Similar to the linear SVM classification, the decision boundary formula in higher-dimensional space uses converted feature vectors to produce the weight vector and bias term. The decision boundary can be stated as follows for non-linear SVM classification:

$$W^{T} \Phi(x) + b = 0$$

Where the input vector x is transformed into a higher-dimensional space by the feature mapping function (x). Using the training examples and the optimization problem, the weight vector w and bias term b are determined.

K- Nearest Neighbors

A non-parametric machine learning approach called KNN (k-nearest neighbors) is implemented for classification and regression. It operates by locating the k training dataset data points that are closest to the new input data point and then assigning a class or value based on the vote of the majority or the average of the k neighbors. The KNN classification formula is as follows:

$$y = model (y1, y2,, yk)$$

Where y is a new data point's projected class label and y1,y2,...,yk are the class labels of the new data points k nearest neighbor's in the training dataset. The mode function gives the k closest neighbor's most prevalent class label. The straightforward formula for KNN in regression is:

$$y = (1/k) * \sum yi$$

Where yi is the value of the new data points ith nearest neighbor in the training dataset and y is the prediction value for a new data point. The formula calculates the values of the k closest neighbors averages.

Logistic Regression

A statistical technique known as logistic regression is used for binary classification, which means it forecasts the likelihood that an input will belong to one of two possible classes. It is frequently utilized in a variety of industries, including marketing, banking, and medicine. Using function to predict the likelihood that an input belongs to one of two groups is the fundamental concept underpinning logistic regression. Any input value can be translated into a value between 0 to 1 using the logistic function, which represents the chance of being in the positive class. The logistic regression equation is:

$$p(y=1|x) = 1/(1 + exp(-z))$$

Where the possibility that the input x belongs to the positive class is given by p(y=1|x). With weights chosen during training, z is a linear mixture of the input features: $z = B0 + B1 \times 1$, $B2 \times 2$, + $Bn \times xn$. The coefficients or weights of the logistic regression model are b0, b1, b2,, and bn. The features are represented by x1, x2,..., and xn. X exp represents the exponential function. To increase the likelihood of the observed data, the logistic regression model finds the optimal values for the coefficients orr weights (b0, b1, b2,, bn) during training. After the model has been trained, the logistic function with the input features may be used to forecast the likelihood that new input would fall into the positive class. The input is categorized as either belonging to the positive class or the negative class depending on whether the probability exceeds a predetermined threshold, which is typically 0.5.

Naïve Bayes

A probabilistic machine learning approach called Naïve Bayes theorem, it makes the assumption that given the class label, the input features are conditionally independent. It is frequently employed in text classification, spam filtering, and sentiment analysis despite its basic premise, and frequently works well in actual application. The Naïve Bayes formula is:

$$P(y \mid x) = P(x \mid y) * p(y)/p(x)$$

When p(y|x) is the high probability that the input x falls under the category y. the conditional probability that input x would be observed given class y is p(x|y). p(x) is the likelihood of witnessing input x, and p(y) is the prior





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probability of class y. The Naïve Bayes algorithm is quick and effective, and it performs well with short training datasets and high-dimensional inputs. The conditional independences assumption, however, could not always be accurate, resulting in subpar performance.

Decision Tree Algorithm

The decision tree is a technique to machine learning that is frequently used for applications in classification and regression. The way it works is to divide the input space into smaller chunks based on the values of the input features, then assign a class or value to each section. Each node in the algorithm represents a test on a certain property, and each branch represents one or more possible test results. This creates a tree-like structure.

Partitioning the input space into areas that are as uniform in terms of class labels as possible is the aim of classification. The method is repeated recursively for each group of data until a stopping requirement is satisfied. The algorithm chooses the best feature to split the data depending on a metric like entropy or the Gini index. The objective of regression is to divide the input space into areas that minimize mean squared error or another regression metric. To reduce error, the method chooses the optimal feature and split point, then iteratively repeats the procedure fir each group of data until a stopping requirement is satisfied.

Random Forest algorithm

Random forest is a supervised learning strategy for problems involving classification and regression. It is an ensemble method that builds several decision trees and combines their predictions to produce a model that is more accurate and reliable. The core idea behind the random forest algorithm is to build a lot of decision trees at random, combine their predictions, and reduce. The steps in the random forest method are as follows: Decide how many trees (n) you want to see in the forest. Choose a portion of the training data and a subset of the features at random for each tree. Create the decision tree using the subset of the data and features that you have chosen. Till all of the trees are grown, repeat steps and 3. Pass a new data point through all trees in the forest to predict it, then take the guess with the most votes. The following can be used to forecast the results of a random forest model:

$\hat{y} = model(f_1(x), f_2(x),...,f_n(x))$

Where \$haty\$ is the expected result, \$f i(x)\$ is the decision trees \$i\$-th output for the input \$x\$, and \$model\$ is the function that gives the decision trees most common output. Instead of the mode, the predicted outcomes in regression problem may be the mean of all the trees output.

XGBoost Classifier

The gradient boosted trees approach is well-liked and well implemented in the open-source framework XGBoost. A supervised learning approach called gradient boosting combines the estimates of several weaker, simpler models in an effort to forecast a target variable with a degree of accuracy. A scalable, distributed gradient-boosted decision tree (GBDT) machine learning framework is called XGBoost, or Extreme Gradient Boosting. For regression, classification, and ranking issues, it is the top machine learning library and offers parallel tree boosting.

Performance Metrics on ML Algorithms

This section assesses and contrasts the algorithms using performance measures like recall, accuracy, and precision. The predictive performance of the model is evaluated using the Cross-Validation technique to estimate how each model performs outside the sample to a fresh dataset that is also known as test data. The reason for using cross-validation approaches is that we fit models to training datasets when we do model fitting. We only have data on how the models perform on training data without cross-validation. In a perfect world, we would test how well the models perform with fresh data in terms of how accurate their forecasts are. In science, the predictive power of a theory is evaluated. It is a popular strategy because it is easy to grasp and produces a less biased or pessimistic assessment of the model skill than other approaches, including a straightforward train/test split.



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Accuracy

To assess how well the classification are performing, accuracy is examined. By dividing the total number of occurrences from the dataset by the ratio of correctly predicted instances, accuracy is calculated. According to mathematics, the classification's accuracy is defined as follow:

Accuracy =
$$\frac{x}{n}$$
 x 100%

Where x represents how many instances cases were successfully categorized and n represents how many instances cases there were in total. Accuracy in binary classification can be determined using the following positives and negatives:

Accuracy =
$$\frac{TN+TP}{TN+TP+FN+FP}$$
 X 100%

where $TN = True\ Negatives$, $TP = True\ Positives$, $FN = False\ Negatives$, and $FP = False\ Positives$ are all possible outcomes. The accuracy of each model on the training dataset, which is divided into two segments (70% for training and 30% for testing On the testing data, following table lists the 7 algorithms' anticipated accuracy. XGBoost has the highest prediction accuracy with 89%. As a result, other performance metrics like recall and precision are also assessed in order to assess and compare the performance of the models.

Precision

Precision is defined as the ratio of true positive results to the total number of anticipated positive observations. It can be stated mathematically as:

Precision =
$$\frac{TP}{TP + TF} \times 100\%$$

The values for each of the seven classifiers precision are displayed in following Table.

Recall

The proportion of accurate positive outcomes to all observations made in the actual class is known as recall, which is also known as sensitivity. It can be written mathematically as follows:

$$Recall = \frac{TP}{TP + FN} \times 100\%$$

The seven classifiers recall values are displayed in Table.

F1 Score

The evaluation matrix known as the F1-score takes the harmonic mean of the two matrices, Precision and Recall, to create a single metric. To put it simply, the weighted average mean of Precision and Recall is the f1 score. In natural language processing, it is also utilized

F1 Score =
$$\frac{TP}{TP + \frac{1}{2}(FP + FN)}$$

The calculated metrics have shown in figure 1, It shows that the Decision Tree and XGBoost classifiers performs best in terms of recall, accuracy, and precision. These findings suggest that XGBoost, out of the seven classifiers ML algorithms, is the most accurate at determining student performance.

CONCLUSION

Machine learning techniques have been extensively used in a number of applications, and they have also been carefully evaluated as useful tool for building knowledgeable machines that can advise academicians on how to make decisions based on Student's background information, performance data in school and colleges. With the survey findings and the data analysis with classification algorithms taken into for predicting the students performance The performance of the existing models, including SVM, Logistic regression, KNN, DT, RFA, Naïve Bayes and XGBoost are compared as a benchmark. Decision Tree and XGBoost outperforms with other algorithms in terms of accuracy on the training and test set. after estimate, the predictive models performance is





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assessed in terms of accuracy, precision, and recall on unobserved data using the k-fold cross-validation procedure to gauge their aptitude. The study's findings demonstrated that the XGBoost Classifier model performs best in terms of accuracy, precision, and recall, F1 scoring correspondingly.

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Table 1. Summary of related work on student performance prediction with various Machine Learning Algorithm.

Related Papers	Features	Methodologies	Dataset Source	Result
Predicting student	Seven publicly	Hybrid Regression	Public datasets.	The research
performance and	available datasets	Model (HRM)-	rublic datasets.	establishes a potent





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its influential factors using hybrid regression and multi-label classification [15].	with various attributes.	1.Collaborative filtering technique 2.Fuzzy rules 3. Lasso linear regression Multi-Label Classification For Predicting Key Factors.		hybrid approach for accurately predicting student academic grades and identifying pivotal performance factors.
Machine learning methods in predicting the student academic motivation [21].	Variables: Assign view(V1), Forum view discussion(V2), Questionnaire view(V3), Resource view(V4).	Neural Networks, Decision trees, and Support Vector Machines.	Faculty of Education in Osijek.	The results showed that the RBF neural network model performed better with classification accuracy of 76.92%.
Classification and prediction of student performance data using various machine learning algorithms [22].	18 Attributes	SCM, Naïve Bayes, C-4.5, ID3	UCI Machinery Student Performance (649 instances)	SVM is better
Predicting student performance using machine learning to enhance the quality assurance of online training via Maharat platform [23].	Gender, Academic Specialization, Academic degree, The number of training courses you attended through the "Maharat" platform.	K-nearest Neighbourhood, Naïve Bayes, Decision Tree, Support Vector Machine.	Maharat platform at Taif University based on the online learning training standards in the Kingdom of Saudi Arabia (KSA).	SVM is better with 93.2% accuracy.
Students performance prediction using KNN and Naïve Bayesian [24]	DOB, Gender, City, Secondary School Name, Specialization, Fathers Job, Student Status	KNN, Naïve Bayes.	Gaza Strip 2015 (500 instances)	Naïve Bayes is better than KNN
Machine Learning Algorithm for Student's Performance Prediction [25].	Class Test Marks, Attendance Mark, Presentation Mark, Assignment Marks, Midterm Marks And Final Examination Marks	Linear Discriminant Analysis, Gradient Boosting Classifier, Random Forest Classifier, SVC, KNN and Decision Tree Classifier.	Daffodil International University Dhaka, Bangladesh. (1170 Student's data)	The most perfect result and accuracy with K-Nearest Neighbors, Decision Tree Classifier model with an accuracy of 89.74% & 94.44%.





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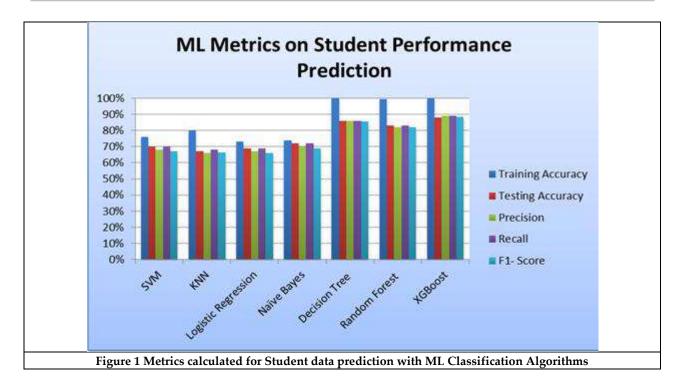
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Student performance analysis system (SPAS) [26].	Quizzes, Assignments, projects, Final examination Grades, Gender, Programs	J-48, Simple CART, BFTree, Random Tree, J-48 Garft	TMC1013 System Analysis and Design in University of Malaysia Sarawak	BF-Tree is better, TMC-1013 System Analysis assist to predict the student performance accurately
Predicting Students' Performance Using Machine Learning Techniques [27].	The dataset contains 161 student records, 76 male and 85 female. The dataset contains twenty attributes.	Artificial Neural Network,Decision Tree, Logistic Regression, Naïve Bayes.	Archeology and the Sociology department of the college of Humanities at Al- Muthanna University	The most accurate model is built using Artificial Neural Network which has an accuracy of 77.04.
Analysis and Prediction of Student Academic Performance Using Machine Learning [28].	22 Attributes.	1. Linear Regression for Supervised Learning 2. Linear Regression for Deep Learning	Kaggle Dataset (648 instances)	Liner Regression for Supervised learning is best.
Student Academic Performance Prediction Using Machine Learning [29]	Name, Grade Id, Semester, Parent Survey, Label.	SVM, Decision Tree, Logistic Regression	Public Dataset.	Highest accuracy for decision tree algorithm which is 96.96.
Prediction of Students Performance using Educational Data Mining [30].	19 Attributes	Naïve Bayes	Amrita School of Arts & Science, Mysure	Naïve Bayes is used for knowledge classification.
A Comparative Study to Predict Student's Performance Using Educational Data Mining Techniques [31].	MEDU, Attendance, 1st semester GPA	Naïve Bayes, Decision Tree	Industrial Engineering University, Islam, Indonesia	Naïve Bayes is performing better.
Using Machine Learning models to predict student retention: Building A State-wide early warning system [32].	15 Attributes	LR, NB, DT	Administrative student data (179517 records)	Predict at risk students with 80% probability.





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RESEARCH ARTICLE

A Stacking Ensemble of Deep Cluster (En-MHSAMKFC-CNN) Based Deep Learning for Cancer Classification Approach

K.Prema*

Associate Professor, Computer Science, Sri Ramakrishna College of Arts & Science (Autonomous), Coimbatore, Tamil Nadu, India.

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*Address for Correspondence

K.Prema

Associate Professor, Computer Science, Sri Ramakrishna College of Arts & Science (Autonomous), Coimbatore, Tamil Nadu, India. Email: prema.k@srcas.ac.in



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ABSTRACT

The method of increasing classifier performance by integrating the contributions of trained sub-models to tackle the same classification issue is known as Ensemble Learning (EL). Overall, each base learner votes, and the meta-learner, which is a model that learns to correct the predictions of the base-learners, gains the final prediction. The ensemble technique outperforms single learners in prediction accuracy. The Meta model receives the output of the sub-models (base learners) as input and then learns to merge the input predictions to produce a final prediction that is better than each of the base-classifiers. The concept of ensemble learning module is structured and named as Ensemble of MHSAMKFC based CNN (En-MHSAMKFC).

Keywords: Ensemble, stacking, backing, over-fitting, under-fitting.

INTRODUCTION

Ensemble methods are strategies for enhancing model accuracy by mixing numerous models rather of utilizing a single model. The integrated models considerably improve the accuracy of the results. It has increased the prominence of ensemble approaches in machine learning.

Categories of Ensemble Methods

Ensemble methods are classified into two types: sequential ensemble techniques and parallel ensemble approaches.





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- Sequential ensemble approaches, such as Adaptive Boosting, create base learners in a sequential order (AdaBoost). The successive production of basic learners fosters dependability among the base learners. The model's performance is then enhanced by giving bigger weights to previously misrepresented learners.
- Parallel ensemble techniques, base learners are generated in a parallel format, e.g., random forest. Parallel
 methods utilize the parallel generation of base learners to encourage independence between the base learners.
 The independence of base learners significantly reduces the error due to the application of averages.

Advanced Ensemble Techniques

Some of the Advanced Ensemble Techniques are listed below

Stacking Method

The ensemble approach is stacking, which is also known as stacked generalization. This approach works by allowing a training algorithm (such as Decision Tree, KNN, or SVM) to combine the predictions of numerous other comparable learning algorithms. Stacking has been used successfully in regression, density estimation, distance learning, and classification. It may also be used to calculate the mistake rate while bagging. This model is used to predict results from the test set.

Bagging Method

Bootstrap aggregating is commonly used in classification and regression, and also known as bagging. Through the use of decision trees, it improves model accuracy while substantially lowering variance. Bagging is classified into two types, i.e., Bootstrapping and aggregation.

Boosting

Boosting is an ensemble strategy that uses prior predictor failures to improve future predictions. The strategy merges numerous weak base learners into a single strong learner, considerably boosting model predictability. Boosting works by placing weak learners in a sequential order, so that weak learners can learn from the next learner in the sequences which result in improved prediction models. Gradient boosting, Adaptive Boosting (AdaBoost), and Extreme Gradient Boosting (XGBoost) are all examples of boosting.

- 1. AdaBoost employs weak learners in the form of decision trees, which typically feature one split, also known as decision stumps. AdaBoost's primary decision stump is made up of observations with equal weights.
- Gradient boosting adds predictors to the ensemble incrementally, with prior predictors correcting their successors, enhancing the model's accuracy. New predictors are fitted to compensate for the consequences of prior prediction mistakes. The gradient of descent assists the gradient booster in identifying and correcting errors in learners' predictions.
- 3. XGBoost makes use of decision trees with boosted gradient, providing improved speed and performance. It relies heavily on the computational speed and the performance of the target model. Model training should follow a sequence, thus making the implementation of gradient boosted machines slow.

ENSEMBLE MODEL APPLICATIONS TOWARDS GEM DATASET

Ensemble approach is an effective strategy for cancer diagnosis that is increasingly being used to integrate various learning algorithms to increase overall prediction accuracy (Dietterich 2000). Much research has demonstrated the promise of ensemble learning for enhancing data classification accuracy under uncertainty (Dietterich 2000; Yang et al. 2010). However, it is both a necessary and sufficient condition for an ensemble to outperform its individual members the base classifiers should be accurate and diverse. AdaBoost, Bagging, Random Subspace, Random Forest, and Rotation Forest are some examples of these approaches (Zhang & Zhang 2008). AdaBoost has grown in popularity because to its ease of use and versatility (Pratama et al. 2018). This approach constructs an ensemble of classifiers by applying a given base learning algorithm to successively acquired training sets produced by either resampling from the original training set or reweighting the original training set based on a set of weights preserved across the training set (Zhang & Zhang 2010). As a result, AdaBoost seeks to generate new "strong" classifiers capable





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of better predicting challenging occurrences for earlier ensemble "weak" members. Wu (2018) suggested a Multi-label Super Leaner based on Heterogeneous Ensembles to increase multi-class super learner classification accuracy. A multi-label classification issue is one in which each sample can represent more than one class label at the same time, for example, an image may be assigned sea and beach or sea and mountains depending on the things it includes. The purview of bioinformatics, yet it was demonstrated to outperform all other approaches for music sentiment analysis, bird acoustics, and scenic datasets. Again, the variety of challenges it handles demonstrates the ability of heterogeneous ensemble to transcend NFL constraints. Yu et al. (2019) proposed a novel method for accurately predicting diagnostically complex cases of cancer patients by combining medical imaging, advanced ML algorithms, and heterogeneous ensembles. They also utilized this technique to demonstrate why some imaging aspects make it difficult to identify even with standard Computer-Aided Diagnosis (CAD) software. To train heterogeneous base classifiers and to construct an ensemble of trained classifiers increased total prediction accuracy to 88.90%, compared to the best accuracy recorded in the literature of 81.17 percent. When compared to other proposed ensemble approaches, such as AdaBoost (Tharwat 2018), Bagging (Jafarzadeh et al. 2021), and Random Forest (Cutler et al. 2011), Rotation Forest is more robust because it can always improve both the generalization ability of individual classifiers and the ensemble diversity. Zhang and Zhang (2008) presented RotBoost, a new ensemble classifier generation approach that combines Rotation Forest and AdaBoost. AdaBoost replaces the basis classifier in the Rotation Forest algorithm in this new ensemble technique. When several non-microarray gene-related data sets from the UCI library are used, the experimental findings reveal that RotBoost outperforms Rotation Forest and AdaBoost.

PROPOSED METHODOLOGY

To handle the over-fitting and under-fitting problems of CNN classifiers, a stacked ensemble model is presented that uses many learning models to build one optimal prediction model. In this model, Stacking based Ensemble model has been used to ensemble the proposed MHSAMKFC-CNN. Every classifier vote is merged to predict the final class label in the stacking ensemble model. As a result, the classifier developed using ensemble learning approaches outperforms the single classifier.

Algorithm 1: Stacking Ensemble Algorithm

```
Input: Data set D = \{(x_1, y_1), (x_2, y_2), ..., (x_m, y_m)\}; First-level learning algorithms L_1, ..., L_T Second-level learning algorithm L.
```

Process

```
1. For t=1,\ldots,T: %Train a first-level learner by applying the 2. h_t=L_T(D);% first-level learning algorithm L_T 3. end 4. 4. D^*=\emptyset; % Generate a new data set 5. For i=1,\ldots,m: 6. For t=1,\ldots,T: 7. z_{it}=h_t(x_i); 8. end 9. D^*=D^*\cup((z_{i1},\ldots,z_{iT}),y_i); 10. end 11. h^*=L(D^*); % Train the second-level learner h^* by applying % the second-level learning algorithm L to the % new data set D^* Output: H(x)=h^*(h_1(x),\ldots,h_T(x))
```





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Working Module of Stacking ensemble with MHSAMKFC-CNN

The construction of proposed stacking ensemble with MHSAMKFC-CNN is given in Figure.1. In this module, stacking learning is employed with MHSAMKFC-CNN classifier to ensemble the features to obtain the final class prediction with majority voting is provided in Figure.2.

Steps Involved in Majority Voting Ensemble process

In this process, the classifiers from the first layer (i.e., training layer) will return the probability of belonging to a class (selected features from MHSAMKFC). In the second layer i.e., testing layer, same MHSAMKFC features are consider as the input of CNN classifier. The classifiers from the first layer will return the probabilities that go into the next classifier input which also utilizes MHSAMKFC of parameters and features. Finally, the output of the classifier are processed to majority votingand to final class predictions (ensemble performance).

IMPROVING THE STACKING ENSEMBLE WITH MHSAMKFC-CNN

The construction of the improved Stacking Ensemble with MHSAMKFC-CNN is illustrated in Figure.3. In this module, the developed ensemble model works by combining FS process (MHSAMKFC) and classification (CNN) process. Combining the models created by each CNN Classifier yields the ultimate best classification performance. The performance for improving the Stacking Ensemble with MHSAMKFC-CNN. An enhanced stacking ensemble model is suggested to properly categorize the GEM dataset for cancer prediction. This model consists of a two-stage method. The basic learners (MHSAMKFC - CNN (1,2...N)) are utilized to create the classifier outputs in the first step. MHSAMKFC - CNN model will boost the variety and robustness of the stacking model, can learn and understand training data from many angles.

RESULT AND DISCUSSION

The effectiveness of existing methods like CPEM (Lee et al. 2019) G-Forest (Abdulla & Khasawneh 2020) and aEKNN and proposed En - MHSAMKFC- CNN are compared based on above mentioned datasets Accuracy, Precision, Specificity, Sensitivity, and F1-Score are the metrics used to evaluate the performance.

Accuracy

The definition of the accuracy values of existing and proposed methods are compared and depicted in Table 3.1. Figure.4 displays the Accuracy of existing CPEM, G-Forest and aEKNN with proposed En-MHSAMKFC-CNN. In this analysis, En-MHSAMKFC-CNN method is 6.39%, 4.21% and 1.65% for Leukemia dataset; 6.75%, 4.03% and 2.56% for Lymphoma dataset and 7.09%, 4.25% and 2.31% for Prostate dataset is higher than that of CPEM, G-Forest and aEKNN methods respectively on given dataset. This analysis shows that the En - MHSAMKFC - CNN can achieve better accuracy than other methods for microarray cancer classification.

Precision

The definition of precision values of existing and proposed methods are compared and depicted in Table 3.2. Figure.5 displays the precision of existing CPEM, G-Forest and aEKNN with proposed En - MHSAMKFC- CNN techniques. In this analysis, En - MHSAMKFC-CNN method is 5.8%, 3.39% and 1.36% for Leukemia dataset; 7.05%, 4.32% and 2.91% for Lymphoma dataset and 6.97%, 4.91% and 2.86% for Prostate dataset is higher than that of CPEM, G-Forest and aEKNN methods respectively on given dataset. This analysis shows that the En - MHSAMKFC - CNN can achieve better precision than other methods for microarray cancer classification.

Specificity

The definition of Specificity values of existing and proposed methods are compared and depicted in Table 3.3. Figure.6 displays the specificity of existing CPEM, G-Forest and aEKNN methods with proposed En-MHSAMKFC–CNN techniques. In this analysis, En – MHSAMKFC - CNN method is 6.78%, 4.59% and 2.02% for Leukemia dataset; 6.28%, 2.94% and 1.75% for Lymphoma dataset, 7.31%, 4.61% and 2.18% for Prostate dataset is higher than





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that of CPEM, G-Forest and aEKNN methods respectively on given dataset. This analysis shows that the En -MHSAMKFC - CNN can achieve better specificity than other methods for microarray cancer classification.

Sensitivity

The definition of Sensitivity values of existing and proposed methods are compared and depicted in Table 3.4. Figure.7 displays the sensitivity of existing CPEM, G-Forest and aEKNN with proposed En-MHSAMKFC-CNN techniques. In this analysis, En-MHSAMKFC-CNN method is 5.43%, 3.72% and 1.85% for Leukemia dataset; 7.27%, 4.53% and 2.11% for Lymphoma dataset, 7.22%, 4.78% and 2.26% for Prostate dataset is higher than that of CPEM, G-Forest and aEKNN methods respectively on given dataset. This analysis shows that the En - MHSAMKFC - CNN can achieve better sensitivity than other methods for microarray cancer classification.

F1-Score

The definition of F1-Score values of existing and proposed methods are compared and depicted in Table 3.5. Figure.8 displays the F1- score of existing CPEM, G-Forest and aEKNN with proposed En-MHSAMKFC-CNN techniques. In this analysis, En-MHSAMKFC-CNN method is 6.27%, 4.64% and 2.00% for Leukemia dataset; 6.98%, 4.93% and 2.68% for Lymphoma dataset, 7.22%, 4.85% and 2.53% for Prostate dataset is higher than that of CPEM, G-Forest and aEKNN methods respectively on given dataset. En – MHSAMKFC – CNN can achieve better F1- Score than other methods for microarray cancer classification.

CONCLUSION

En-MHSAMKFC- CNN is suggested for development to overcome the over-fitting and under-fitting problem caused by imbalanced class label in micro array gene datasets for cancer classification. En-MHSAMKFC- CNN is compared with different existing methods in terms of Accuracy, Precision, Specificity, Sensitivity and F1-Score using three microarray datasets such as Leukemia, Lymphoma and Prostate micro array datasets. The efficiency of these strategies is examined in MATLAB 2018a, which runs on Microsoft Windows 7 with a 2.70 GHz Intel CPU and 4GB RAM. Finally, simulation results show that En-MHSAMKFC- CNN is more effective than traditional classifiers at improving and quantifying the performance of cancer prediction systems that are used to improve and quantify cancer types GEM datasets, balancing performance efficiency with less computational complexity.

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Table 1 Evaluation of Accuracy

Datasets\ Classifiers	Accuracy				
	CPEM	En- MHSAMKFC- CNN			
Leukemia	90.55	92.44	94.77	96.34	
Lymphoma	91.38	93.77	95.11	97.55	
Prostate	92.05	94.56	96.35	98.58	

Table 2 Evaluation of Precision

Tubic 2 Livaruation	Table 2 Evaluation of Trecision						
Datasets\ Classifiers		Precision					
	СРЕМ	G- Forest	aEKNN	En- MHSAMKFC- CNN			
Leukemia	92.66	94.87	96.77	98.09			
Lymphoma	91.43	93.82	95.11	97.88			
Prostate	92.53	94.34	96.22	98.98			

Table 3 Evaluation of Specificity

Datasets\ Classifiers	Specificity						
	CPEM	CPEM G-Forest aEKNN En-MHSAMKFC-C					
Leukemia	91.32	93.24	95.58	97.52			
Lymphoma	92.01	94.99	96.10	97.79			
Prostate	92.14	94.52	96.77	98.88			

Table 4 Evaluation of Sensitivity

3							
Datasets\ Classifiers		Sensitivity					
	CPEM	CPEM G-Forest aEKNN En-MHSAMKFC- C					
Leukemia	93.87	95.45	97.21	99.01			
Lymphoma	92.23	94.65	96.89	98.94			
Prostate	92.31	94.46	96.79	98.98			

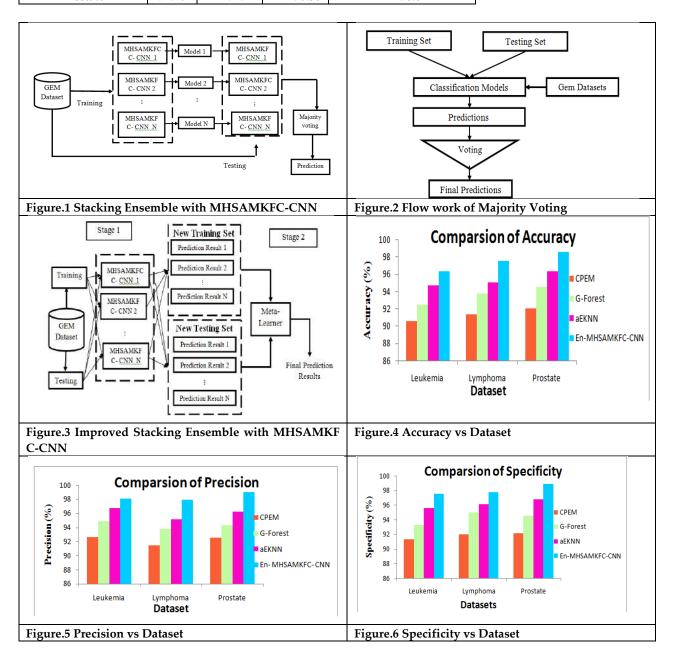




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Table 5 Evaluation of F1- Score

Datasets\ Classifiers	F1- score			
	CPEM	G-Forest	aEKNN	En- MHSAMKFC- CNN
Leukemia	92.92	94.37	96.81	98.75
Lymphoma	93.37	95.19	97.28	99.89
Prostate	92.16	94.24	96.38	98.82

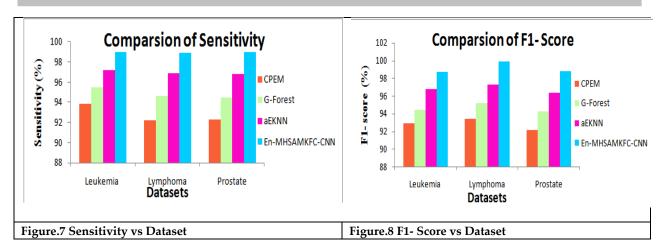






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REVIEW ARTICLE

A Comphrensive Review and Meta-Analysis on Psychological Distress among Students Using Machine Learning Techniques

Maria Suman A¹ and Saleema J S^{2*}

¹Researech Scholar Computer Science, CHRIST (Deemed to be University) Bengaluru, Karnataka, India. ²Associate Professor, Statistics and Data Science, CHRIST (Deemed to be University) Bengaluru, Karnataka, India.

Received: 30 Dec 2023 Revised: 29 Jan 2024 Accepted: 13 Mar 2024

*Address for Correspondence Saleema J S

Associate Professor, Statistics and Data Science, CHRIST (Deemed to be University) Bengaluru, Karnataka, India. Email: saleema.js@christuniversity.in

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ABSTRACT

The COVID-19 pandemic poses a risk to students' physical and mental well-being. This descriptive crosssectional study is currently assessing the mental health of children and adolescents, and it is evident that their well-being is being significantly affected by the high rates of depression, stress, and anxiety that are being reported. The situation is further compounded by home confinement, which raises concerns about the negative impact of screen usage on their1mental health. Thus, ameta-analysis is currently being conducted to examine the global psychological1distress rates during the1COVID-19 pandemic. Additionally, the1study includes various moderator analyses to explore factors contributing to the variability of depression anxiety and stress prevalence rates. Studies were1selected based on inclusion1and exclusion1criteria obtained from PubMed, Scopus, Google Scholar, Web of Science, MedLine, and Nature following the PRISMA model. The pooled prevalence rate of depression was 35% (95% CI,-0.35-0.38;), the anxiety was 22% (95%CI, 0.40-0.44;); stress was 28% (95%CI, 1.56-2.24;)However, the outcomes differ based on age, circumstances, and location. Moreover, the sample size of these studies was small. The random forest outperformed and is considered suitable machine learning for data analysis and study. The latest research highlights how important it is to give teenagers access to 1 mental health1services both during and after the pandemic in order to make sure they get the help they need.

Keywords: Depression, Anxiety, Stress, Students, Machine Learning





Maria Suman and Saleema

INTRODUCTION

Globally, the multitude of people spending time at home has sharply increased since the outbreak of Coronavirus-19 in early 2020. In addition, a by-product of the Coronavirus disease is an intense feeling of stress, fear, and anxiety that occurs in such an uncertain environment. Reports from the World Health Organization (WHO) reveal that even before the pandemic, 1 in 3 people suffers from anxiety and depression. The study looks at five important life areas that could have affected adolescents during the pandemic: expectations for the future, physical activity and media exposure, social interactions, mental wellness, and conduct difficulties. As far as adolescent social relationships are concerned, the pandemic has substantially changed their social lives due to infection control measures such as the closure of schools and the physical separation of students from each other [2]. This social deprivation may disrupt peer relationships. Adolescents are commonly concerned about being disconnected from friends due to the pandemic . Additionally, there's a chance that the pandemic harmed family dynamics: in the event of a pandemic, parent-child relationships may suffer due to stress from confinement, caring responsibilities, and financial instability[1]. On the other hand, adjustments to family attention and routines as well as more time spent together may have benefitted certain children [3]. Globally, mental illness has reached epidemic proportions, and WHO estimates that one-fourth of the population will suffer from a mental or neurological disorder sometime in their lives. Mental illness is a result of imbalances in brain chemistry. During the pandemic, children and adolescents experienced various psychosocial and behavioral problems, including lack of attention, hesitation, disinterest, and fear of asking questions about the pandemic. The current pandemic has long-term health effects as it is a lingering stressor; assessing their mental wellness and suggesting appropriate treatments is essential. Adolescents frequently struggle with mental health issues like stress, anxiety, and depression.

Depression symptoms include sadness, fear, uncontrollable worry, aggression, and disruptions in sleep and appetite. When researchers observe heterogeneity across studies, as during COVID-19, it is often a sign that demographic, geographical, and methodological moderators are necessary to understand the relationship, it is possible to detect who is more likely to experience higher prevalence compared to lower prevalence, based on moderator analyses. Before and amidst the COVID-19 pandemic, prevalence rates of mental illness differ based on child age by experiencing higher rates of internalizing disorders. It is also possible that methodological characteristics of studies influence estimations of prevalence rates. Study methods of poorer quality may overestimate prevalence rates, for instance. To determine possible prevalence rates, it is also necessary to consider the timing of data collection. The early months of the pandemic may have been more stressful and overwhelming than later, but school closures and extended social isolation may have likely exerted psychological strain on some students. Machine learning and psychology come together whenever the research question is "How capable is it to predict?" With the advancement of technology, mental health researchers and doctors can collect more information quicker than ever before. Smartphones, neuroim aging, and wearable technologies have made this possible. The application of machine learning for the study of these data has improved reliability over time. The aim of machine learning is to develop computers that are able to learn on their own by applying advanced probabilistic and statistical techniques.

Objective

To understand about mental wellness is crucial, considering the global epidemic proportions of mental illness and the lingering stressor effect of the pandemic, which can lead to various psychosocial and behavioral problems among students. Thus the current systematic review and meta-analysis aim to (1) Identify adolescents' mental health problems amidst the pandemic. (2) Illuminate machine learning strategies to improve mental health issues for students.





Maria Suman and Saleema

Study Methods Search Strategy

This systematic review followed Preferred Reporting Items for Systematic Reviews and Metaanalyses (PRISMA)reporting guidelines. The literature review focused on the Web of Science (WoS) database, Scopus, MeSh (MedLine and Google Scholar) Keywords, NATURE, and PubMed database. A search strategy was devised by incorporating three central themes: mental illness (including depression, stress, and anxiety), children and adolescents (age 10 to 18 years), the COVID-19 pandemic, and Machine Learning. Unpublished pre-prints were also searched.

Selection Criteria (Inclusion and Exclusion Criteria)

In terms of the selection criteria, the following criteria were set: (1) articles specifically targeting children up to 18 years of age; (2) papers that relate COVID-19 impact on adolescents and children; and (3) studies analyzing the psychological behavior of confined children (4) machine learning techniques that support in the prediction of mental illness. (5) The study taken was Empirical Articles written in English were considered in the study. The titles and summaries of each article were read first to determine whether they met the criteria mentioned above, which allowed us to rule out papers that did not. In the end, nine scientific papers were selected after a more thorough reading of the articles chosen (Figure 1)

Data Extraction

The following coding processes were used in order to extract the data from the articles: (1) author(s) and year of publication; (2) research title; (3) place/country of publishing; and (4) key concepts of the research. Prevalence rates for anxiety, stress, and depression were derived from study publications that satisfied inclusion requirements.

LITERATURE REVIEW

Elizabeth A. K. Jones et al.[19] aims to assess how the pandemic affects adolescents' mental health. In this pandemic, stress, extended home confinement, worry, and social media overuse can negatively impact adolescents' mental health. COVID-19 poses mental challenges to adolescents around the world. Adolescents require physical and psychological support to grow, develop, and thrive, regardless of the uncertainty caused by the current crisis. It is important to make use of all therapy modalities and resources available to assist adolescents in adjusting to the pandemic's effects. In the aftermath of COVID-19 and similar disasters, there needs to be more research on improving adolescent mental health. To better address adolescents' psychological needs, it may be suggested that telemedicine be implemented globally, among many other interventions. Ray, Suman, et al.[8]study focuses on Coronavirus infection can affect a person of any age, but children are particularly vulnerable because of the disease's complex psychological effects. During the pandemic, certain children would have been put through to quarantine restrictions. In addition, the lockdown has also been reported to have affected children's mental health due to greater access to digital devices. The COVID-19 pandemic and lockdown have also hurt children's mental health. Several questionnaires, such as Short Self-Rating Questionnaires (SSRIs), were used to assess the level of stress experienced by children and youth between the ages of 9 and 18 in the aftermath of the COVID-19 epidemic. The study was cross-sectional and observational research using an online survey to collect data. 369 school kids took part. The stress levels were classified into Low, Moderate, and Severe groups using a Score Scale and analysis to categorize them. In analyzing the total score levels of the Delhi+Mathura zone (n=369), 30.08% (n=111) were found to be under a moderate stress level, 62.87% (n=232) under a severe stress level and 7.08% (n=26) were found to be under an extreme stress level. In Delhi and Mathura combined, a significant difference was found in stress levels between males and females (p*0.04). Delhi and Mathura zones did not significantly differ in stress levels. Children and adolescents must receive proper stress management given the current environment. Family and child needs may be met with an intervention strategy based on culturally appropriate and empirically supported interventions. Kumar, Abhinit, et al.[7]studies concern that mental illness in children and adolescents rises when they transition from childhood to adolescence as the conditions become more complex and intense. This study aims to assess and





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compare anxiety and depression prevalence among suburban and rural children in eastern Uttar Pradesh and to understand their socioeconomic impact. The 11- to 18-year-old age children were divided into two groups; 100 from rural Tikri and 100 from suburban Sunderpur. They collected information about their sociodemo graphics. Screening for depression and anxiety in children was conducted using the Children's Depression Inventory and the Revised Children's Manifest Anxiety Scale. The present state examination was performed using the International Classification of Mental and Behavioural Disorders as the basis for the final diagnosis. A Chi-square test was used to analyze the data statistically. Depressive disorders are prevalent in 14.5% of the population, while anxiety disorders are prevalent in 15%. Rural and suburban areas were not significantly different (P > 0.05) regarding depression or anxiety. Among adolescents, females, and people in the lower-middle economic bracket, depression, and anxiety were more prevalent than in other age groups. Students in classes 9th - 12th were more likely to suffer from depression, while students in lower classes were more likely to experience anxiety. There was a higher prevalence of depression in joint families. The differences between these two groups indicate some crucial trends relating to the factors that impact these problems. Jeelani, Asif, et al.[9]paper is based on during the Coronavirus pandemic of 2019, routine activities were disrupted severely, significantly more deaths were reported, and more morbidity has reported. The mental health needs of adolescents are exceptionally high.

Adolescents attending school in the Kashmir valley of India were evaluated for the prevalence and determinants of depression and anxiety. In January and February 2021, school-going adolescents aged between 15 and 19 years were screened for depression and anxiety using the Patient Health Questionnaire for Adolescents and Generalised Anxiety Disorder questionnaire. A total of 426 (97.03%) out of 439 adolescents responded to the survey. With a mean age of 17.5 + 1.26 years and 57% males, the adolescents were in the middle of their adolescent years. Approximately 16% of people with depression had a history of COVID-19 infection in the past. Twenty percent of adolescents had anxiety. Boys were affected at a rate of 14% and girls at a rate of 27.5%. A significant public health problem is associated with anxiety and depression among adolescents. Based on our study, it appears that adolescents who struggle with self-esteem need immediate assistance. Ms. Sumathi et al.[10] study focuses on treating mental health problems effectively; early diagnosis is crucial. The earlier it is treated, the greater the chance of improving the quality of life for patients. Therefore, it is imperative to address fundamental mental health problems among children that threaten their mental health. If not treated early, these conditions can lead to more complex issues. Machine learning approaches currently well serve in medical data analysis and diagnosis. Five fundamental mental health problems have been diagnosed by comparing eight machine learning techniques on different accuracy measures. To train and test the performance of the methods, a data set of sixty cases is collected. There are 25 attributes identified in the documents that are important for diagnosing the problem. A Feature Selection algorithm has been applied to the complete attribute data set to reduce the attributes. The accuracy of various machine learning approaches has been compared on a complete attribute set and a target attribute set. There is little difference between their performance over the complete attribute set and the selected attribute set among the three classifiers: Multiclass Classifier, Multilayer Perceptron, and LAD Tree.

Usman, Muhammad, et al.[11] study deals with mental disorders such as depression that negatively affect a person's mental health, causing them to behave, think, and feel negatively, leading to physical or emotional problems. In such situations, people lose hope for the positive and only see the negatives. In Machine Learning (ML), a model can be trained on training datasets to predict whether someone will develop depression in the future. Researchers have previously used machine learning techniques to predict depression. A study reviewed ML techniques' effectiveness in predicting depression in older adults. This study has used a variety of classifiers, including Bayes Nets, logistic regressions, multilayer perceptrons, sequential minimizations, decision tables, and random forests. Large datasets can achieve maximum accuracy and precision of 89% and 0.95, respectively. Ahuja, Ravinder, and Alisha Banga [12] study analyzes college students' stress experiences at various stages of their lives. Examination pressure and recruitment stress affect students in often go unnoticed. The dataset was sourced from the Jaypee Institute of Information Technology and contained 206 students' data. We will analyze these factors' impact on a student's mind and correlate this stress with internet usage. As performance parameters, sensitivity, specificity, and accuracy are measured for these four classification algorithms: Linear Regression, Naive Bayes, and SAnd and





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Random Forest Using 10-fold cross-validation further enhances data accuracy and performance. Support Vector Machines were the most accurate (85.71%). Haque, Umme Marzia, et al.[13] goal of the study is (1) to create an algorithm that can predict depression in adolescents and children between the ages of 4 and 17, (2) to evaluate the results to determine which algorithm performs best, and (3) to relate it to family activities and socioeconomic challenges. This study draws data from the YMM, the second Australian Child and Adolescent Survey of Mental Health and Wellbeing 2013-14. A low correlation between the variable of yes/no value and the target variable (depression status) has been eliminated. To detect depression among highly correlated variables with the target variable, we used the Boruta algorithm in conjunction with a Random Forest (RF) classifier. The Tree-based Pipeline Optimization Tool (TPOT classifier) was used to choose appropriate supervised learning models. Depression detection was conducted using the use of RF, Decision Tree (DT), XG Boost (XGB), and Gaussian Naive Bayes (Gaussian NB). Even with differences in performance between models, RF could predict depressed classes 99% with 95 % accuracy, 99% precisely, and in 315 milliseconds (ms) faster than all other algorithms. As well as outperforming all four confusion matrix performance measures and execution duration, this model is more accurate in predicting child and adolescent depression.

Qasrawi, Radwan, et al.[14] aims to identify depression and anxiety risk factors in school children using machine learning. The study included 5685 students studying in public and refugee schools in the West Bank in grades 5-9, aged 10-15. Data were collected using the health behaviors questionnaire for school children during the 2013-2014 academic year, and machine learning was used to predict risk factors associated with mental health symptoms in students. The prediction techniques used were a Decision Tree, Support Vector Machine, Neural Network, Random Forest, and Naive Bayes. For depression and anxiety, the SVM and Random Forest models were the most accurate (SVM=92.5%, RF=76.4%; SVM=92.4%, RF=78.6%). As a result, the SVM and Random Forest produced the best classification and prediction results for the depression and anxiety of the students Shailly Gupta et al.[15] focused on Indian students who were assessed for psychological and physical health due to increased screen usage during COVID-19. With the help of self-made questionnaires and snowball sampling, a cross-sectional survey was carried out on Indian students. The survey received 210 responses from students at various schools and colleges between 15th August and 30th August 2020. The relationship between screen time and physical and psychological health during COVID-19 was statistically significant (t= 19.96; p=0.01). As a result, students should be incorporated into appropriate preventive health measures.

To obtain more reliable and accurate results, further research is needed to identify the health effects of the overuse of screens in specific populations such as adolescents and adults. Twenge, Jean M., and W. Keith Campbell, et al.[16] study focused on an array of psychological well-being measures. Comprehensive screen time standards (including electronic games, cell phones, electronic devices, computers, and TV) were assessed in a large (n = 40,337) national random sample of children and adolescents in the U.S. in 2016. In studies involving daily screen time for 1 hour or more, reduced psychological well-being was associated with a reduction in curiosity, a decrease in self-control, a greater tendency to distractibility, a lower sense of emotional stability, and difficulties with friendship. The risk of ever having been diagnosed with anxiety (RR 2.26, CI 1.58, 3.22), depression (RR 2.39, 95% CI 1.54, 3.70), or treated by a professional in mental health (RR 2.22, CI 1.62, 3.03) or with medication treated for a psychological or behavioral issue was twice as high among 14- to 17-year-olds who were high screen users (7+ hours per day compared to 1 h/day) in the past year. Moderate screen use (4 hours/day) also negatively impacted mental well-being. There was no significant difference in well-being between low users and non-users of screens. Adolescents are more likely to have lower psychological well-being from screen time than younger children.

RESULTS

A total of 100 nonduplicate records were found in our electronic search. To evaluate all 65 full-text articles against inclusion criteria, 65 abstracts were reviewed, and 15 non-overlapping studies met all the requirements.





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Study Characteristics

A total of 15 studies were included in the survey literature and 5 in meta-analyses, of which five studies were on depression, anxiety, and stress. Four studies had depression and anxiety,3 had stress-related issues, and three papers related screen time and depression, anxiety, and stress. The mean (SD) percentage of depression was 22.7%, anxiety was 68%, stress was 25%, and the mean age was 13.0 years (range, 8-19 years).

Pooled Prevalence of Depressive, Anxiety, and Stress Symptoms

The pooled prevalence from a random-effects meta-analysis of 5 studies revealed a pooled prevalence rate of depression is 0.35 (95% CI,-0.35-0.38;) or 35%. The between-study heterogeneity statistic was 0.02. The pooled prevalence rate of anxiety is 0.22 (95%CI, 0.40-0.44;) or 22%, and the heterogeneity statistic was -1.16. Lastly, the pooled prevalence rate of stress is 0.28 (95%CI, 1.56-2024;) or 28%, and the heterogeneity statistic was 1.01. The funnel plot was symmetrical (Figure 2, Figure 3, and Figure 4 in the supplement).

Performance Metrics of Machine Learning Methods

At the citation screening stage of a systematic review of depression, anxiety, and stress the ML approaches were applied. By measuring sensitivity, specificity, accuracy, and AUC (Area under Curve) on a validation set of papers were able to assess the performance of the ML approaches. Different algorithms like Random Forest, Support Vector Machine, Naïve Bayes, Artificial Neural Networks, Decision Tree, XGB, and Gaussian NB were considered to find the best approach for the prediction of depression, anxiety, and stress. The comparative study on algorithms states that Random Forest outperforms other methods by acquiring 1.00 sensitivity,0.85 specificities,0.99 Accuracy, and 0.86 AUC.

DISCUSSION

The COVID-19 pandemic and its associated restrictions and consequences have significantly affected adolescents' psychological well-being. Among other factors, social isolation, peer interactions, and reduced contact with buffering supports (e.g., teachers, and coaches) may have contributed to these increases.4 Additionally, school-based services are the primary source for mental health treatment for 80% of children. The closure of these facilities has rendered these services unavailable to many children. An increase in depression and anxiety rates was observed as the month of data collection progressed. A cumulative association could be formed between social isolation, financial difficulties, missed milestones, and school disruptions. This possibility is still lacking longitudinal research, which is urgently needed. A further longitudinal study on Children's health will be required to confirm the long-term mental health effects of the pandemic on youth mental illness as the pandemic progresses and during the recovery phase. Study results showed that all psychological interventions analyzed benefited healthy and mentally ill adolescents and children. Thus, schools become a suitable environment for facilitating community involvement and family, and they, therefore, have the potential to enhance positive relationships between teachers and students, parents and teachers, and students and their peers. It is important that parents and teachers are involved in the teaching and learning process. Families with no training in this area might have some difficulties with these active methods that are utilized in schools. [20]. The results of emerging research42 indicate that routines reduce child depression and conduct problems during COVID-19. Children and families can mitigate the effects of COVID-19 on youth by implementing consistent and predictable practices around sleep, schoolwork, screen use, and physical activity. Children experiencing clinically elevated mental distress should be provided with additional resources and referred to clinical professionals. As a result, school closures and recreational activities should be considered a last resort. Further, there should be an emphasis on the equitable distribution of mental health resources to children while increasing scalability.





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LIMITATIONS

A systematic review has been conducted in which a set of interventions has been assessed for adolescents and children without explicitly distinguishing between them. There is a limitation here since adolescents and children might need different levels of support regarding their psychological and behavioral health. There may be differences in outcomes that have not been considered in this review. The current meta-analysis includes a reexamination of cross-regional differences will be important once additional data from underrepresented countries become available once the recent meta-analysis includes child and adolescent mental illness estimates across regions. Secondly, most studies were cross-sectional, which precluded long-term research on COVID-19's effect on child mental health.

CONCLUSION

The meta-analysis results show that children's anxiety and depression symptoms were elevated during the COVID-19 pandemic. It also stresses the importance of interventions and recovery efforts to improve the well-being of students. Still, it also emphasizes that when focusing on interventions, individual differences must be considered (e.g., gender, age, COVID-19 exposure). It was found that computer/internet, mobile phone, and television/videogame use were both associated with negative outcomes. Long-term research on this epidemic is necessary to enhance understanding of the implications of the COVID-19 pandemic on today's children's mental health trajectory. This should include studies using pre- and post-COVID-19 measurements. Researchers have found that machine learning can help understand psychiatric disorders through existing studies and research. For machine learning models to perform well, the data samples and their characteristics must be considered. Preprocessing activities like data cleaning and parameter tuning can also influence machine learning models. Observing and classifying mental health problems at an early stage can assist in further treatment in the future.

AREA OF CONFLICT

Behave of all authors the corresponding author states that there is no area of conflict

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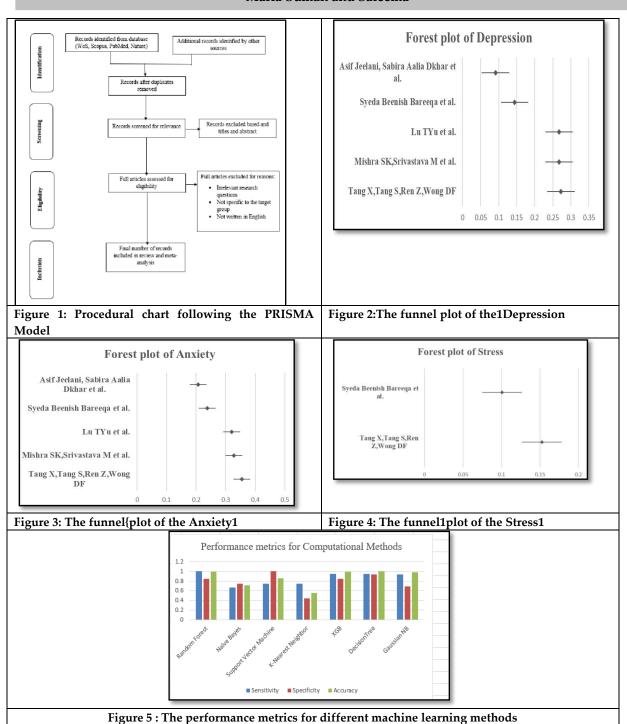
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REVIEW ARTICLE

Determinants of Maternal Mental Health In India: A Systematic Review and Meta- Analysis

Anugraha Mathew^{1*} and J S Saleema²

¹Post Graduate Student, Department of Statistics, and Data Science, CHRIST (Deemed to be University) Bengaluru, Karnataka, India.

²Associate Professor, Department of Statistics and Data Science, CHRIST (Deemed to be University) Bengaluru, Karnataka, India.

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*Address for Correspondence

Anugraha Mathew

Post Graduate Student, Department of Statistics and Data Science,

CHRIST (Deemed to be University) Bengaluru, Karnataka, India.

Email: anugraha.mathew@stat.christuniversity.in



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ABSTRACT

Maternal mental health is an essential component of public health, and it poses a multifaceted challenge in India. Despite advances in maternal care, there is rising concern about the prevalence and consequences of common mental health disorders among pregnant women. Through addressing existing knowledge gaps and investigating barriers related to the recognition and treatment of these disorders, this research strives to provide insights that can guide the formulation of interventions and policies tailored to the cultural nuances of the population. This study aims to conduct a systematic review and meta-analysis to identify the determinants of common maternal mental disorders in India. While targeting the specific objectives outlined in Sustainable Development Goal 3 (SDG 3) concerning mental health and well-being, this study extends its focus to include a broader spectrum of determinants. This approach recognizes the interconnected nature of maternal mental health with various societal factors, including those related to SDGs addressing poverty, hunger, inequality, education, and economic growth. We implemented the PRISMA model for the systematic review, performed a meta-analysis using the random effects model and assessed the publication bias. A pooled prevalence rate of 44.9% (95% C.I = 34.4 to 56) was observed, with a high level of heterogeneity among the studies. In light of the study's findings, policymakers may consider prioritizing interventions addressing the identified determinants that not only improve mental health outcomes but also contribute to broader societal well-being. The results of this study motivate an expanded review of relevant literature to enhance the robustness and generalizability.





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Keywords: Maternal Mental Health, Common Mental Disorders, Systematic Review, Meta-Analysis, Sustainable Development Goals (SDGs)

INTRODUCTION

According to WHO (World Health Organization), maternal health is women's health throughout pregnancy, childbirth and the postnatal period. Over the years, the primacy of maternal health has increased, and India has considered it an essential attribute for development with respect to increasing equity and decreasing poverty. Maternal mortality and morbidity are significant concerns for investing in maternal health. Sustainable Development Goals (SDGs) 2030 has incorporated the aim of reducing the global maternal mortality ratio (MMR) to under 70 per 100,000 live births and ensuring universal access to healthcare services, including family planning, education and information services 1. In association with UNICEF and UNFPA, WHO is determining best practices for providing evidence-based, integrated, people-oriented maternal and perinatal healthcare services. It also aims to provide information and proposals on care models at various phases of the obstetric transition across countries. Many studies have been performed to know the number of deaths, attributes affecting them and their influence on maternal health, maternal morbidity and mortality. Statistics suggest that for every maternal death, around 20 - 30 women suffer from maternal morbidity. Pregnancy is always characterized by joy, enthusiasm and psychological well-being, but it is a common misconception that mental health stays strong throughout pregnancy and the postpartum period. This has led to around 80% of unreported maternal mental health conditions 1. Maternal mental health conditions extend beyond depression, anxiety, and psychosis, including issues such as baby blues, maternal dysthymia, OCD, birthrelated PTSD, and others. Globally, around 10% of pregnant women and 13% of postpartum women encounter a mental disorder, predominantly depression. These figures are notably higher in developing nations, reaching 15.6% during pregnancy and 19.8% following childbirth3.

In India, based on a systematic review and meta-analysis published in the Lancet Psychiatry in 2020, the prevalence of perinatal depression was 22.9%, and the prevalence of anxiety was 15.7%. There are certain demographic segments that face an elevated risk of mental disorders due to increased exposure and susceptibility to adverse social, economic, and environmental conditions, closely tied to gender-related factors. Prevalent among these mental disorders are common mental disorders (CMDs), primarily encompassing conditions such as depression and anxiety disorders. This research seeks to fill knowledge gaps and investigate hindrances to the identification and treatment of maternal CMDs, with the ultimate goal of guiding the creation of culturally sensitive interventions and policies. Additionally, it strives to foster maternal mental well-being, strengthen the bond between mothers and infants, and enhance the long-term health outcomes for mothers and their children in India. The objective of this paper is to synthesize available evidence and generate collective estimates regarding the prevalence of maternal common mental disorders (CMDs) in India. This information aims to guide the development, implementation, and assessment of interventions aligned with the National Health Mission (NHM) goals. Our approach involves identifying evidence and conducting a meta-analysis based on existing studies that investigate the prevalence of CMDs in India. Furthermore, we intend to compile an overview of the determinants influencing maternal CMDs across the country. It emphasizes the criticality of addressing interconnected societal factors such as poverty, hunger, inequality, educational access, and economic development, as outlined by SDGs, to promote optimal maternal mental health outcomes. This review will particularly focus on the prevalence rates of depression and anxiety, representing the two most commonly observed CMDs globally.

METHODS

Data Source and Search Strategy

A comprehensive systematic review identified the pertinent evidence for maternal mental health in India. The Preferred Reporting Items for Systematic Reviews and Meta-Analyses (PRISMA) 2020 guidelines were followed for the review. Preceding the initiation of the review, we established a meticulously formulated research question, a





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well-defined search strategy, and explicit inclusion-exclusion criteria. The review utilized PubMed and Google Scholar as the primary databases and also extracted papers from different web sources. We assessed over fifty-eight articles published from the year 2013 till July 2023 to guarantee alignment with the current prevalence and factors influencing maternal CMDs. The systematic review was conducted through two approaches: firstly, systematic searches of databases were performed, encompassing all sources relevant to maternal common mental health in India and grey literature. Secondly, hand searches involved scouring reference lists of cited systematic reviews for additional relevant studies. We incorporated the MeSH terms and free-text word synonyms for "Maternal Mental Health", "India", "Common Mental Disorders", "Depression", "Anxiety", "Perinatal", "Antenatal" and "Postpartum". The review was undertaken on an expedited schedule to guide research priorities before the inception of the project proposed at NHM. Consequently, it was impractical to scrutinize all the results from the databases or acquire inaccessible papers.

Inclusion-Exclusion Criteria

We took into account the research studies across diverse geographical locations in India. These papers revolve around the investigation of maternal mental health and factors associated with it, particularly within the context of the perinatal period. The inclusion criteria encompass the examination of incidence, prevalence, and risk factors associated with maternal common mental disorders (CMDs) such as postpartum depression, antenatal psychiatric symptoms, and stress during the perinatal and postpartum period. The study designs include both community-based and hospital-based cross-sectional and longitudinal approaches, contributing to a thorough investigation of maternal mental health and its determinants across diverse settings and populations in India. We rejected studies based on the exclusion criteria listed: studies using sample sizes of less than 20 people; data gathered from countries other than India; study methodologies other than observational, such as literature reviews and qualitative investigations; studies where the data could not be extracted separately; studies that report on the prevalence or determinants of CMD without using a recognised diagnostic tool or screening tool; and full-text papers that were not retrievable during the review period.

Data Extraction and Screening Strategy

We imported and organized the search outcomes from all the sources using End Note software, eliminating any duplicate entries in the article list. Subsequently, we conducted screening based on the titles and abstracts of the articles, leading to the retrieval of relevant papers. Due to time limitations, papers that could not be retrieved were excluded. Finally, we thoroughly reviewed the full text of the selected papers and retained those that fulfilled the eligibility criteria. The extracted data from the eligible papers were stored in Microsoft Excel which included the first author's name, the title of the paper, the type of study, year of publication, data source, the type of population, the state of India, sample size, event of interest, event size, prevalence of CMDs, the type of maternal common mental disorder, its measurement scale, and its key determinants.

Statistical Analysis

In light of the expected heterogeneity in outcomes among the included studies, attributed to the limited number of extracted studies, we deemed a meta-analysis investigating explanatory factors of heterogeneity to be more valuable than a systematic review devoid of quantitative results. It could be argued that, with the acknowledgement of limitations in synthesizing heterogeneous results, quantitative syntheses offer richer insights compared to a qualitative interpretation of results and allow exploring diversity in outcomes. We performed a random effect meta-analysis in Comprehensive Meta Analysis (CMA) software. The forest plots were used to evaluate the pooled estimate of the prevalence of maternal common mental disorders in India. Cochran's Q-statistic tests whether there is a significant difference in the common effect sizes among all the studies obtained after the systematic review due to sampling error alone. However, Q statistic depends highly on the size of the meta-analysis, hence relying on statistical power so we suggest not to infer results based on it. Furthermore, we calculate I², in simple terms, the percentage of variability not caused by sampling error, which is highly dependent on the precision of the included





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studies. Therefore, we infer meta-analysis results based on its I² value and 95% prediction interval. Also, a funnel plot is used to study the publication bias in the studies.

RESULTS AND DISCUSSION

Study Description

The study outlines the process of selecting reviews in 0. We acquired a total of 58 records considered to be potentially relevant, consisting of 50 records identified through database searches and 8 from alternative sources such as grey literature. Among the 58 records, 8 duplicates indexed in more than one database were excluded. Subsequently, the remaining 50 records underwent initial screening based on their titles and abstracts, excluding 2 records that did not meet the inclusion criteria. Full-text reviews were conducted on the remaining 42 reviews as we excluded 6 papers that were not retrievable, revealing 6 studies that satisfied the inclusion criteria. Expanding on Naaz et al.'s discoveries, our systematic review aligns with their observation that women in lower socioeconomic classes, with less education and more children, are more likely to experience common maternal mental disorders (CMD)[4]. Furthermore, Naaz et al.'s subsequent study emphasizes the significance of advanced maternal age, nuclear family structure, urban living, and single parenthood as notable factors contributing to maternal CMDs[5]. To deepen our understanding, George et al.'s research suggests that socioeconomic status, recent adverse events, perceived support, and maternal health concerns are crucial determinants of postpartum depression[6]. Bhaskar, echoing Naaz, identifies a higher prevalence of postpartum depression in urban Indore [7], while Sheeba et al.'s study in Bangalore reveals a notable connection between domestic violence, pregnancy-related anxiety, and prenatal depression[8]. Expanding the scope, Vijayaselvi et al.'s work emphasizes the importance of pregnancy planning and the husband's employment status as key factors linked to maternal stress[9]. Our systematic review aligns with Edge's study on correlates of perinatal depression, affirming the significance of factors such as smoking, gestational diabetes, and pre-eclampsia in influencing maternal mental health outcomes[10]. In most of these 6 papers, they assessed the effect of sociodemographic variables, obstetric history, and postpartum factors on mental health disorders. The most common factors affecting mental health were social status, parity of mothers, domestic violence, recent catastrophic events, and the intentionality of the pregnancy. Due to the variability and inconsistency of factors identified in the literature review, we are impeded from constructing a meta-analysis for these variables.

Meta-Analysis

We conducted a random model meta-analysis on the six studies that satisfied our inclusion criteria and obtained six estimates on CMDs. The pooled prevalence estimate of maternal CMD is 44.9% (95% C.I. = 34.4 to 56, I^2 = 93%). Here, we observe that a high variance of 93% in observed prevalence reflects variance in true prevalence rather than sampling error. We also obtained that the true prevalence rate is not the same for at least two of the studies in the analysis using Cochran's Q test (Q = 69.591 with 5 degrees of freedom (d.f.) for α = 0.05). 0depicts the forest plot of prevalence rates of maternal CMDs in India. Despite the limited size of our meta-analysis, we established 95% prediction intervals for the prevalence rates, ranging from 14.2 to 80.1, providing a glimpse into the nature of predicted intervals in this study. With an expanded analysis size, the interval length is expected to significantly decrease, yielding a more precise estimated range.

Limitations

In prevalence studies that aggregate data from diverse observational studies involving patients with varied characteristics and conducted in various settings, there is anticipated to be heterogeneity in the estimates. Nevertheless, despite our endeavours to execute a systematic review of high quality, it is probable that the quality and heterogeneity of the included studies influence the reliability of the findings. While we investigated variations in estimates based on factors such as the use of diagnostic interviews or screening tools for identifying depression and/or anxiety, the diversity in screening tool types, translation and validation processes, and variations in how and by whom screening was conducted might account for some of the identified heterogeneity. It is also conceivable that





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there are relevant studies we may have overlooked because of time constraints. The assumption of the generalizability of the term "Maternal Common Mental Disorders" (CMD) might also have introduced disparities in the estimates.

CONCLUSION

Despite limitations in prevalence estimates, this study reveals a high burden of maternal CMDs in India, associated with socio-demographic factors, obstetric history, and domestic violence, rather than solely pregnancy-related causes. The alarming prevalence of 44.9% reinforces the urgent need for multi-pronged policy interventions aligned with India's SDG 2030 goal for mental health. Implementing policies like micro-loan programs, skills training, and childcare support for low-income mothers could potentially alleviate financial burdens and improve mental health outcomes. In addition to that, establishing community-based support groups and peer mentoring programs could address social isolation and empower mothers with knowledge, potentially reducing depressive symptoms. While further research is needed to optimize interventions and address diverse contexts, these initial steps hold immense potential to strengthen maternal mental health and build a more equitable and supportive society.

Future Scope

The systematic review has the potential to be revised by incorporating the remaining papers from the databases, facilitating a comprehensive meta-analysis encompassing all available records to date. Additionally, there is an opportunity to perform subgroup analyses, providing a more nuanced understanding of the prevalence of maternal Common Mental Disorders (CMDs) among various subgroups within the existing literature in India.

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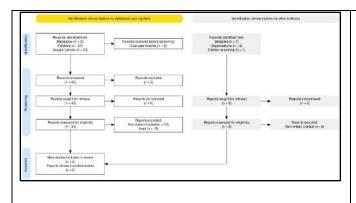
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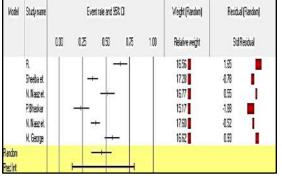


Fig 1: PRISMA diagram of the selection process

Fig 2: Forest plot of prevalence rates





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RESEARCH ARTICLE

Enhanced Method of Emotion Detection and Face Recognition for **Attendance Monitoring Using Machine Learning**

Jeyalakshmi^{1*}, S. Deepak Prakash² and N. Sanjay Kumar³

¹Associate Professor, Department of Information Technology, Sri Ramakrishna College of Arts & Science, Coimbatore, Tamil Nadu, India.

²PG Scholar, Department of Information, Technology, Sri Ramakrishna College of Arts & Science, Coimbatore, Tamil Nadu, India.

³PG Scholar, Department of Information Technology, Sri Ramakrishna College of Arts & Science, Coimbatore, Tamil Nadu, India.

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*Address for Correspondence Jeyalakshmi

Associate Professor, Department of Information Technology, Sri Ramakrishna College of Arts & Science, Coimbatore, Tamil Nadu, India. Email: jeyalakshmi@srcas.ac.in



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ABSTRACT

The use of facial recognition is essential in every industry in the modern era. One of the most popular biometrics is face recognition. Along with many other benefits, it can be used for security, authentication, and identity. Additionally, facial recognition software can be used to record attendance in work environments, institutions, and colleges, among other places as well. The sustainable attendance system seeks to streamline the attendance management process and save time and effort for both administrators and attendees. Automated facial recognition technology eliminates the need for manual data entry and reduces the administrative burden associated with attendance tracking. It allows administrators to allocate resources more efficiently and focus on more value-added tasks. The currently available manual method takes a lot of time and work to maintain, so the goal of this system is to build up a facial recognition-based class attendance system. Moreover, there may be opportunities for proxy participation. The machine learning project, titled " Emotion Detection and Face Recognition for Attendance Monitoring" trains a model to identify student attendance based on their emotions during class using Python, OpenCV, and student datasets. Facial emotion recognition and classification are done using the CNN and Haar Cascade algorithms. It ensures accurate attendance records by capturing and analyzing facial features using computer vision algorithms. The goal of this project is to offer an alternative to the manual attendance-taking approach that is currently in use by staff members.





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Keywords: Face Recognition, Face Detection, Emotion Detection, Haar Cascade Classifier, CNN, Open-cv

INTRODUCTION

In many schools and universities, the traditional technique of recording attendance is a laborious effort. The faculty members who must manually call out each student's name to record attendance are further burdened, as this could take up to five minutes for the entire class [1]. To do this, a lot of time is needed. In certain situations, proxy attendance is permitted. As a response, several institutions started using other techniques to monitor attendance, like the use of fingerprint, iris, and radio frequency identification (RFID) [2]. These methods, however, are obtrusive and queue-based, which could take additional time [14]. Face recognition has developed into an essential biometric characteristic that is non-intrusive and simple to learn. Systems that rely on facial recognition are not very sensitive to different expressions on the face [5]. Face identification and verification are the two categories that make up a face recognition system. [3] While face verification compares a face image to a template face image in a 1:1 matching procedure, face verification compares a query face image to a 1: N problem. The technique of automatically extracting, analyzing, and comprehending meaningful information from a single image or a series of photos is known as open-source computer vision. Colleges could use computer vision to use facial recognition technology to electronically take attendance. This will not only help faculty members save valuable time, but it will also teach staff members and their students how to code and the range of computer vision applications that are feasible [15].

College instructors and staff would have access to more student data and early notice of notable changes in emotional state if there was an emotion detection system in place [8]. While security cameras are deployed in many institutions, they are rarely used for anything other than capturing images. Computer vision can be implemented in college settings with these cameras. Instructors and staff can then track student involvement and emotion using the data gathered. A possible method to evaluate students' mental health might include computer vision [6]. The many applications of open- source computer vision, such as face detection, face identification, and emotion analysis, could be useful in tackling the problem of college students [11]. If a student exhibit repeated signs of depression or anger, our program will be able to alert the guidance counselor at the college in advance. Using the help of these capabilities, educators can use facial recognition to save the time spent on outdated attendance procedures and connect themselves and their students with computer vision and deep learning applications [9]. Facial recognition and emotion detection can help teachers to monitor the changes in their student behaviors to increase productive knowledge among the student and helps in achievement [13].

Facial Recognition Technology

The system utilizes advanced facial recognition algorithms to identify and authenticate individuals based on their unique facial features [4]. This technology analyzes key facial and marks to create a unique biometric template for each person.

Real-Time Facial Detection and Emotion Detection

The system employs computer vision techniques to capture and analyze facial images in real-time [7]. It detects faces within a given frame and matches them against the pre-registered biometric templates to authenticate individuals [12]. Emotions expressed by the students are detected using the Haar Cascade Classifier and CNN.

LITERATURE REVIEW

Face Recognition System Using Deep Learning, Saibal Manna, Sushil Ghildiya, Krishnakumar Bhimani[16]Future expansion of the company will benefit from the face recognition attendance system's real-time video processing design, that supports company growth. The objective of this paper is to develop a real-time video





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processing-based facial recognition attendance and time tracking system. Four research experiments were conducted in this study: the accuracy rate of the face recognition system during real-time register; the stability of the face recognition time and attendance system with real-time video processing; a study of the face recognition attendance system's skip rate using real-time video processing; and a configuration of the face recognition attendance system's interface. The experimental results indicate that the face recognition technology and computer assistance used by the time and attendance system to accomplish the desired time and attendance results fully reflects the feasibility design of the overall algorithm. Students those who are successfully completed the attendance check-in method did so in a timely manner, eliminating the need for the convoluted roll call sign and instantly realizing its purpose. The conversion form for the attendance system and the future system time have achieved significant advancements that have improved the face recognition technology's dependability and attendance rate. Our scientists should investigate and realize it more thoroughly. Face Recognition Attendance System Based on Real Time Video Processing, Xiaofeng Han, Hao Hang, [17] In this article, A face recognition attendance system that utilizes real-time video processing is created, and two institutions within a province are chosen to have their student attendance checked in real-time. This article primarily lays out four approaches to look at the issues: the truancy rate of the face recognition attendance system with real-time video processing, the stability of the face recognition attendance system with realtime video processing, and the accuracy rate of the face recognition system during the actual check-in. Analyzing the facial recognition attendance system's interface settings with real-time video processing is challenging.

Through an analysis of these issues, a face recognition technology-based attendance system concept is put forth, and research is done on face recognition attendance systems based on real-time video processing. According to research findings, the video facial recognition system has an accuracy of roughly 82%. With just roughly 13% of students missing class, the facial recognition time attendance system and manual fingerprint punching are more reliable and accurately identify check-ins. Efficiency is significantly higher than in the control group, which helps stop pupils from departing early and missing lessons. Facial Recognition Using Haar Cascade and LBP Classifiers, Anirudha B Shetty, Bhoomika, [18] Facial Recognition is the biometric technique used in face detection. The facial recognition technology is used to validate or identify a face from the multi-media pictures. Facial recognition has become increasingly crucial as civilization has grown. Face detection and identification have increased globally. It is due to the need for security, including authorization, national security, and other exigent situations. Numerous algorithms exist for the purpose of facial recognition. The goal of this research is to compare two face recognition methods that are standardized for classification: Haar Cascade and Local Binary Pattern. The precision of Haar Cascade surpasses that of the Local Binary Pattern as a result, but its execution time also exceeds it.

Attendance System with Emotion Detection: A case study with CNN and OpenCV, Herman Kandjimi, Abhinav Srivastava, [19] The way multinational corporations (MNCs) care for their workforce is by asking for input, where daily mood is considered a non-factor. Manual data collection techniques are conventional and widely used, and they consume employees' productive time to perform. This procedure can be automated with the use of a certain solution. This study investigates the use of machine learning (ML) and artificial intelligence (AI) models in an automated attendance and mood system. Because of the way the system is set up, businesses can use facial recognition and a customized Convolutional Neural Network (CNN) model to generate data on an employee's mood or emotions. The entrance and exit of each block or floor may have cameras installed by the companies. Cameras are able to take pictures and feed them into OpenCV, making integration simple and attendance tracking possible. The photo that was taken can also be used to identify an employee's mood and be stored in their own database. On the same day, the Human Resources (HR) department can monitor employee facial expressions to get input about new projects, hikes, seminars, and social events.

PROPOSED SYSTEM

Before their photos are taken and added to the dataset, each student in the class must register by providing the necessary information such as their name, department, and regno. Faces from the live streamed classroom video will be identified during each session. A comparison will be made between the faces identified and the photographs





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included in the dataset. In the event a match is made, the relevant student's attendance will be recorded. With Open CV, Attention Detection uses students' facial expressions in a classroom to provide an Excel sheet with the results. The Attention Detection using Face Emotion Recognition machine learning project trains a model to identify student attendance based on their emotions during a class using Python, OpenCV, and a dataset of students. The goal of this project is to offer a substitute for the manual way of taking attendance that is currently used by teachers.

Real-Time Monitoring and Reporting

Real-time tracking and reporting of attendance data is made possible by the system. Administrators have instant access to attendance records, are able to track trends in attendance, and create thorough reports. This makes it possible to make decisions quickly, act quickly when necessary, and handle attendance-related data effectively.

Integration with Existing Systems

Existing staff or student information systems can be easily integrated with the sustainable attendance system. This breaks down data silos, streamlines data administration, and improves administrative effectiveness all around. Integration makes it easier to synchronize attendance data with other pertinent systems, which streamlines procedures all around.

Improved Data Analysis and Insights

For enhanced data analysis and insights, the system's automated data administration and consolidated database are advantageous. Administrators can monitor students' moods through the use of Haar Cascades classifiers, which enables them to make appropriate plans for the following class hour. Facial expression detection is one of the many computer vision applications in which CNNs have demonstrated impressive performance. Using a dataset of labeled facial expressions, we may train a CNN model to identify patterns and characteristics linked to various emotions.

Reduced Administrative Burden

Staff members' administrative workload is lessened with automated attendance tracking since it eliminates the need for human data entry and monitoring, freeing up time for other crucial responsibilities. Typically, this process can be divided into four stages,

Dataset Creation

Images of students are captured using a web cam. Next step is to resize the cropped images to particular pixel position (1280*720) dimension. The dataset used in this project consists of images of students captured during a class hour. Each image is labeled with the corresponding emotion of the student, such as happy, sad, or neutral. The dataset was collected using a camera placed in the classroom. In the dataset the image will contain the name and register number of the students as the name of the image.

Face Detection

Face detection here is performed using Haar-Cascade Classifier with Open source computer vision. Haar Cascade algorithm needs to be trained to detect human faces before it can be used for face detection. Features like edge, line, and center surround are the main frames as shown in the fig.2. This step is called feature extraction. The haar cascade training data used here is in the form of an xml file-haarcascade_frontalface_default. The haar features will be used for feature extraction where the objects in the image are identified as shown in Fig.3.

Face recognition and Emotion detection

The machine learning model used in this project is trained using the labeled dataset. The training process involves training a Convolutional Neural Network (CNN) to recognize emotions in the images. The model is then fine-tuned to detect the level of attention of each student in the class based on their emotions, during recognition process image of the face to be recognized and then compared with the already computed image and returns the best matched label associated with the student it belongs to.





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Attendance Updation

After face recognition and emotion detection process, the recognized faces will be marked as present in the excel sheet and the rest will be marked as absent and the list of student's emotions will be filtered and given to the respective faculties. Faculties will be updated with an hourly attendance sheet at the end of every class.

RESULTS AND DISCUSSIONS

The image of the students is captured in the webcam as shown in fig.4. and stored in the folder named Class_data as shown in fig.5. The system employs computer vision techniques to capture and analyze facial images in real-time. It detects faces within a given frame and matches them against the pre-registered biometric templates to authenticate individuals. The students are supposed to enter all the required details in the student database. Once the class starts, the web cam starts automatically. Camera pops up and starts detecting the faces in the frame as shown in fig.6. Then it automatically starts clicking photos of the students through live streaming camera. These images then will be pre-processed and stored in training images folder. The Fig.6, and Fig.7. shows the face recognition window where two registered students are recognized and if in case, they were not registered it would have shown 'unknown'. The faculties will get the hourly report through excel sheet as shown in fig.8. This is important because the list of presented students with their emotional state during the class will be recorded, it shows the emotional state like Neutral, happy, sad, angry etc. An emotion detection would give college faculty, the more information on students and early warning of significant shifts in emotional state. The staff can identify the student's behavior through the hourly report of attendance with emotion of the students.

Pseudocode- Face recognition in Frame

faces In Frame = face_recognition.face_locations(imgS) encode In Frame= face_recognition. face_encodings (imgS, faces In Frame) for endode Faces, face Loc in zip(encode In Frame, faces In Frame): is matched = face_recognition.compare_faces(all Encodings, endodeFaces) face D is= face_ recognition. face_ distance (all Encodings, endode Faces) try:

best Match Index = numpy.argmin (face D is) except Value Error as ve: print(ve)

CONCLUSION

Every student in the class needs to register by providing the necessary information. After that, their photos will be taken and included to the dataset. Faces will be identified from the classroom's live streaming footage during each session. The identified faces will be contrasted with photos that are included in the collection. Should a match be discovered, the relevant student's attendance will be recorded. The faculty will receive an excel sheet with the hourly report at the conclusion of each session. This system uses facial recognition algorithms to create an efficient way to track students' attendance in class. Face ID will be able to be used by the proposed system to track attendance. Through the use of a webcam, it will first identify faces. Once a student has been identified, their attendance will be noted along with any emotions they have expressed, and the attendance record in the Excel spreadsheet will be updated.

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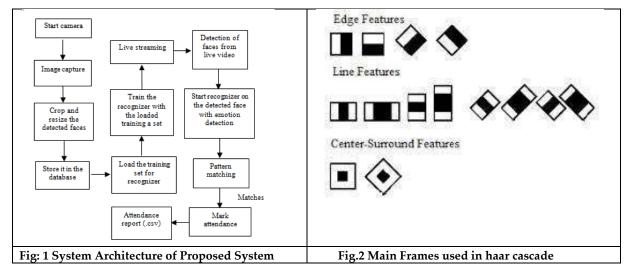
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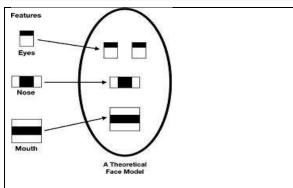




Fig.3. Haar Features

Fig.4. Face Detection



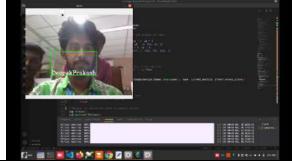


Fig.5. Students image folder



Fig.6. Face Recognition of student 1

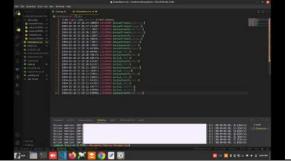


Fig.7. Face Recognition of student 2

Fig.8. Hourly report of attendance with emotion of the students





RESEARCH ARTICLE

Performance Analysis of Gray Level Image Segmentation by Using **Different Methods**

T.Brinthaguru¹ P. Karthick^{2*}, and P.Vinoth³

¹Research Scholar, Department of Mathematics, Kalasalingam Academy of Research and Education, Krishnankoil-626 126, Virudhunagar, Tamil Nadu, India.

²Assistant Professor, Department of Mathematics, Kalasalingam Academy of Research and Education, Krishnankoil-626 126, Virudhunagar, Tamil Nadu, India.

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*Address for Correspondence

P. Karthick

Assistant Professor, Department of Mathematics, Kalasalingam Academy of Research and Education, Krishnankoil-626 126, Virudhunagar, Tamil Nadu, India.

Email: karthickphd91@gmail.com



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ABSTRACT

Image segmentation is of great importance in understanding and analysing objects within images. Using segmentation and subsequent color density evaluation, the procedure entails turning hazy images into meaningful and practical ones. Medical and industrial areas, among others, frequently use the image segmentation method. In this paper, we will discuss four important types of image segmentation methods to segment gray images and compare the accuracy of segmentation based on the error ratio test.

Keywords: Image segmentation, Grayscale image, Edge-based method, Threshold-based method, Region Based method, Clustering based method.

INTRODUCTION

Using a variety of methods, image processing aims to improve images or extract relevant information from them. In order to improve viewer comprehension, this method usually takes an input image and produces either a transformed image or pertinent characteristics connected with it. Automated analysis and interpretation can be facilitated by information extracted from photos. In computer vision, image thresholding is still a major obstacle. Because segmentation problems are ill-posed, getting generalized image segmentation results remains a challenge despite years of focused research. Of all the segmentation techniques available, graph theoretical approaches have shown promise as workable solutions. These methods provide interesting insights into tackling segmentation





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difficulties and increase the value of computations by mathematically modeling image elements into well-defined structures. A key method in digital image processing and analysis is image segmentation, which is breaking a image up into separate areas or sections, frequently according to pixel properties. This could involve grouping related areas according to color or shape characteristics or separating foreground components from the background. Image segmentation is an important process in medical imaging, especially for recognizing and labeling pixels or voxels within a 3D volume that represent anomalies such as brain tumorsor other organs. It is possible to improve the look of these abnormalities for better identification by adjusting pixel densities and contrasts. Of all the segmentation techniques available, thresholding is one of the most straight forward and efficient. Because of its high accuracy and adaptability, it is frequently used for image segmentation jobs. Image processing covers a wide range of applications and standards, and MATLAB is a great interactive tool for working on different image processing problems. The goal of this work is to explore the nuances of image segmentation, with a particular emphasis on thresholding and edge detection techniques. This paper will examine many illustrative examples of image segmentation using grayscale images as a foundation, concluding with an analysis of error ratios. This paper is organized as follows: In section 2, we briefly discuss relevant literature to contextualize our work within the existing research landscape. Section 3 presents our methodology, outlining the segmentation methods employed and discussing their applicability to our study. Following this, Section 4 details the experiments carried out and presents the corresponding results. Finally in section 5, we conclude our work by summarizing key findings and suggesting avenues for future research.

Related Work

In [1] contains A key component of many visual comprehension systems is segmenting images or video frames into discrete objects and segments. The outstanding performance of deep learning-based segmentation algorithms across several image segmentation tasks and benchmarks has attracted a lot of attention. These algorithms' design and methodology allow them to be divided into several architectural categories. Color information and texture features are commonly used in creating histograms or distinguishing borders, limits, and textures inside pixels, as 2012 [3] demonstrated. As such, segmenting data based on color and texture is a widely used strategy for indexing and data management. Abdul-Nasir et al. [2] produced the results of using a novel method for color image segmentation that combines the k-means clustering algorithm with a variety of color models. The research explores the examination of different color components in RGB, HSI, and C-Ycolor models in order to identify the component that produces the best segmentation results. Furthermore, procedures such as median filtering and seeded region growing are a extraction were used to smooth the image and remove unwanted regions, respectively, in order to improve the quality of the image. Kumar et al [10] discussed the RGB model is displayed, along with abreakdown of the RGB image into its R, G, and Bcomponents. We turn a color image into a grayscale image using MATLAB, and then the grayscale image's pixel intensity varies based on the suggested transformation equation. Fan et al. [6] present a revolutionary automatic image segmentation method in 2001.

First, color edges in a image are automatically detected by combining a fast entropic thresholding methodology with an improved isotropic edge detection method. Moreover, color-edge extraction results combined with Seeded Region Growing (SRG) improve image segmentation accuracy. We also investigate the automatic face detection use of this image segmentation system. The technique's usefulness in directed segmentation tasks, such as segmentation-driven image compression techniques and medical image segmentation, was developed by Mohsen et al. [4]. An overview of different image segmentation techniques is given in this publication by Mohsen et al. [5]. These techniques are mostly used for the identification of points, edges, line patterns, and other features. Notably, for applications like pattern recognition and medical imaging, writers have used methods like SegNet, UNet++, and encoding-decoding using A trous. A novel thresholding approach is presented by Narayana moorthy S et al. [8] with the goal of determining the optimal threshold value for grayscale images. It makes use of the Normalized graph cut measure in conjunction with Atanassov's intuitionistic fuzzy sets. In particular, the degree of doubt or proficiency in identifying whether a pixel in the image refers to the backdrop or the item within the image is indicated by Atanassov's intuitionistic fuzzy index values.





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This study [7] uses a normalized graph cut metric as the foundation for thresholding with the goal of separating objects from the background by applying the traditional membership function. The fuzzy normalized graph cut method is the name given to the algorithm suggested in this paper. The suggested approach is analyzed using MATLAB software, and the article concludes with a summary of the findings for further study and recommendations. provide a fuzzy set-based thresholding method in 2023[9]that is easy to use, efficient, and uses a lot less processing power. First, the membership function is used to express the color image. Second, the membership values on the image are generated by integrating the spatial and intensity data to create the symmetrical weight matrix.

Existing Gray Scale Image segmentation Methods

Threshold based segmentation

This technique is used to generate binary images from originals. This is accomplished by comparing pixels to a predefined threshold value. Pixels larger than this value will be assigned 1(black); pixels less than or equal to this value will be assigned 0 (white). The algorithm's steps are as follows:

- 1. As input, read an image.
- 2. Create new 2-D variables and store the image size in them.
- 3. Enter the threshold value.
- 4. Create a new zeros matrix with the same dimensions as the image.
- 5. Match pixels to the threshold value.
- 6. Show the results. If we need to split the items in the new image, we need to write them down.
- Step 1. Take an image as input
- Step 2. Convert this image to grayscale.
- Step 3. Provide the threshold value.
- Step 4. Show only pixels greater than the threshold value.

Edge based segmentation

This method is used to determine the details of objects by using intensity-level (grayscale) detection. Edge detection is one of the most common strategies used to analyse images. The edge method uses the difference between colors to determine the details of the edges. These details appear as white lines on a black background. There are several different methods used to filter

- Sobel filter
- Prewitt filter
- Robert's filter
- Log filter
- Zero-cross filter

This function (Edge function) has two parameters: the image's variable and the method. The steps of the algorithm are as follows:

- 1. Read an image.
- 2. Use edge function and any of the aforementioned filters.
- 3. Display the image.

Watershed segmentation

Watersheds provide a free object-splitting approach that is particularly useful for dividing objects that are in contact with each other. This process understands images as physiological surfaces in which the value of f (x, y) corresponds to elevation. The watershed algorithm finds the watershed basins and hill lines in the image where water would theoretically collect. Suppose that there is a perforated hole at each regional low point and the natural terrain is submerged from below by water rising through these holes at a uniform rate. Pixels are marked as submerged as the





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water rises. Ultimately, water will rise to a level where two submerged areas become integrated. When this happens, the algorithm creates one thick pixel dam separating the two areas. As the flood continues, the entire image is divided into separate collection basins. The steps for this algorithm are as follows:

- Step 1. Read an image.
- Step 2. Pre-process black border removal.
- Step 3. Adapt the watershed area.
- Step 4. Estimate the Lesion ratio.
- Step 5. Merge watershed areas.
- Step 6. Define watershed borders.
- Step 7. Smooth borders (Border sampling and b-spline smoothing).
- Step 8: Overlay borders

Clustering based segmentation

Clustering is an undirected technique used in data mining for identifying several hidden patterns in the data without coming up with any specific hypothesis. The reason behind using clustering is to identify similarities between certain objects and make a group of similar ones. There are two different types of clustering, which are hierarchical and non-hierarchical methods. Non-hierarchical Clustering In this method, the dataset containing N objects is divided into M clusters. In business intelligence, the most widely used non-hierarchical clustering technique is K-means. Hierarchical Clustering In this method, a set of nested clusters are produced. In these nested clusters, every pair of objects is further nested to form a large cluster until only one cluster remains in the end.

K means

K-means clustering is an unsupervised technique no labeled response for the given input data. K-means clustering is a widely approach for clustering. Generally, practitioners begin by learning about the architecture of the dataset. K-means **Clustering Method** clusters data points into unique, non-lapping groupings.

RESULTS AND DISCUSSIONS

Image Segmentation Approaches

A set of images are used to measure the implementation of the suggested algorithm along with some of the frequently used algorithms given in the literature, Performance estimation and comparisons are executed by using real images, where the object can be exactly separate from the background using some appropriate threshold method. The examples are shown in figure 1 and 2. Each figure shown the (a) Original image, (b) Histogram of the Original image, (c) Threshold image, (d) Edge based image, (f) Watershed image, (e) K means Clustering method for image segmentation purpose.

Error Ratio

Error ratio (ER) is the most crucial comparative metric. ER is the proportion of pixels between the object pixels and the optimal threshold image obtained by each method that are placed in the erroneous area. Wrongly segmentation as provided in the following formula,

$$E_R = \frac{N_F + N_{miss}}{100} \times 100$$

 N_{F}

where N, and Nmiss are the number of segmentation pixels in the incorrect area and N, is the number of pixels in the segmented images. In this regard, the value of ER ranges between 0 and 100, with a value 0 denotes a lack of resemblance between the ground truth image and the segmentation result. A value of 100 also indicates a good segmentation result.





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Error Ratio Table

Table 1: Comparison of Error Ratio Result of three Images for Different Image segmentation methods. The original threshold image pixel values and the optimum image pixel values are shown in the Table l. In Table l, the error indicates the numerical values corresponding, in each case, to the similarly among the Edge based method. K means Clustering method and Watershed method. Then we choose from a set of error ratio value the best one. We have noticed in bold type the lowest error value. Please note that in all of the cases. It is noticeable from Table 1 reveals that the different types of image segmentation methods comparison of error ratio.

CONCLUSION

Image segmentation algorithms are capable of transforming opaque images into meaningful ones through which vital information can be extracted, especially in the medical field. This study reviewed some of the hashing algorithms that can be used in MATLAB and looked at the results of each one and the error rate. Finally, we observed that researchers need to improve the low error ratio and reduce computational time in the existing algorithm.

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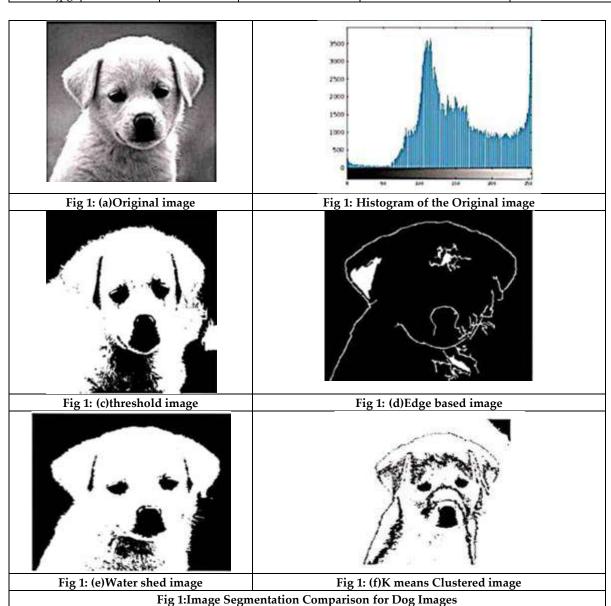




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Table 1: Error Ratio Table

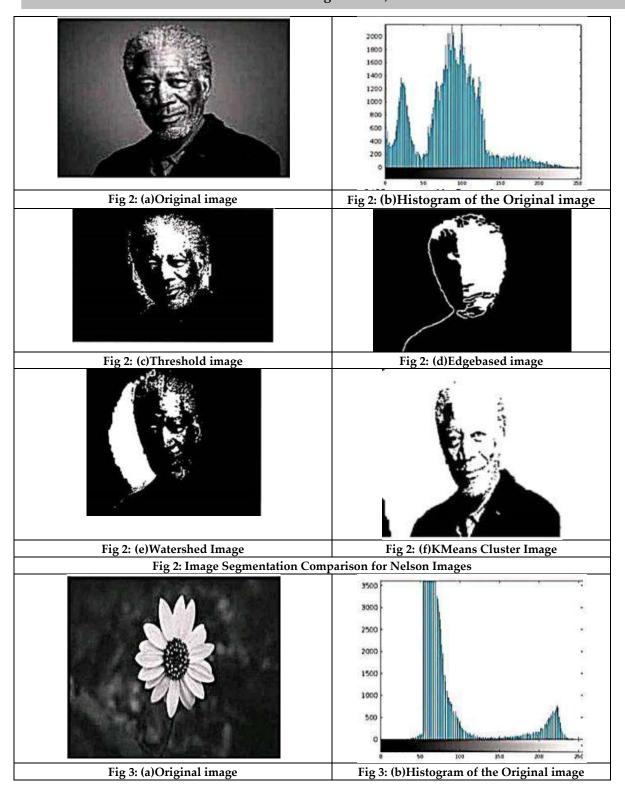
-	Threshold Method		Error Ratio with %		
Images	Original	Optimum	Edge based method	K means Cluster method	Watershed method
	Image Pixel	Image Pixel			
Dog.jpg	298,800	99,600	18.767%	28.28%	15.3%
Nelson.jpg	150,738	50,246	1.07%	8.51%	0.2%
Flower.jpg	150,696	50,232	3.4%	16.5%	2.8%







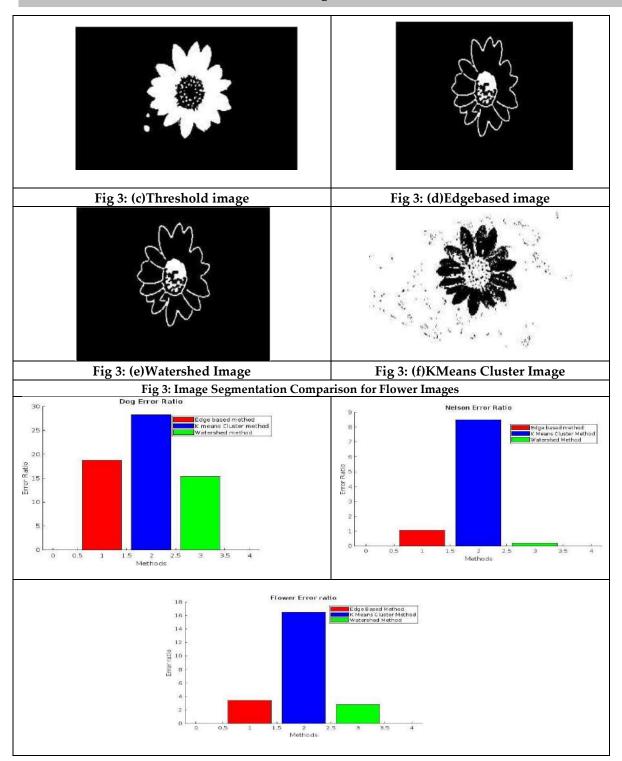
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RESEARCH ARTICLE

A Hybrid CNN, LSTM, and Stacking Ensemble Approach for Improved **Sentiment Analysis**

Kanimozhi.J* and A.Balraj

Assistant Professor, Department of Computer Science, Sri Ramakrishna College of Arts & Science, Coimbatore, Tamil Nadu, India.

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*Address for Correspondence Kanimozhi.J

Assistant Professor, Department of Computer Science, Sri Ramakrishna College of Arts & Science, Coimbatore, Tamil Nadu, India. Email: kanimozhi.j@srcas.ac.in



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ABSTRACT

Opinion mining or Sentiment analysis make use of the enormous volumes of user-generated material that are available on the internet. It is possible to describe Opinion Mining (OM) or Sentiment Analysis (SA) as the process of finding, extracting, and categorizing views on anything. It uses natural language processing (NLP) to track how the public feels about a certain regulation. Making data-driven decisions that affect product development, marketing tactics, and customer service enhancements may be done with this information. To concentrate and classifications feeling from printed information, this examination offered an interesting way to deal with assessment mining that mixes profound learning and AI draws near. Convolutional brain organization (CNN) and long transient memory (LSTM) models are joined in this recommended methodology to increment arrangement exactness. In addition, using the stacking ensemble machine learning method will improve the performance of the suggested strategy even more. The textual material is initially preprocessed in this manner to get rid of extraneous information and noise. Including CNN.

Keywords: Opinion mining, Hybrid Neural Network, CNN, Machine learning

INTRODUCTION

An area of regular language handling (NLP) called assessment mining, ordinarily alluded to as opinion investigation, centers around the PC examination of individuals' verbally expressed sentiments, sentiments, and mentalities [1]. Opinion mining has grown more significant in a range of areas, including marketing, politics, healthcare, and customer service [2] because to the internet's exponential increase in user-generated information.





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Opinion mining aims to identify and extract subjectivity from text, such as reviews of products, comments on social media, and news articles, and to categorise it as good, negative, or neutral [3]. This might offer insightful information on public opinion, assist businesses in improving their goods and services, and have an impact on legislative choices [4]. The primary contributions and objectives of this manuscript may be summarized as follows

- 1. Dataset Pre-processed by using Standard Scalar
- 2. PoS Tag applying using LDA
- 3. Dataset has trained by using Faster R-CNN with Bi-LSTM algorithm
- 4. Feature selection by using Extra tree classifier with Gradient Boosting algorithm
- 5. Opinion Classification using stacking Ensemble Classification

BACKGROUND STUDY

The goal of this research was to give a review and update on the sentimental analysis that is often conducted via social networking. Numerous publications and studies in this sector have been published and completed. Arti, Dubey, K. P., & Agrawal, S. [5] Authorized users may use Twitter's API to gather data and insights from tweets to analyze public opinion for sporting events such as the 2016 Indian Premier League. The result reflects the population's positive and negative opinions. This kind of opinion analysis might give valuable information to the organization and assist them in detecting an unfavourable change in Twitter audience understanding. Early detection of unfavourable trends allows businesses to make educated choices about focusing certain parts of their services and goods on boosting consumer satisfaction. Da'u, A.et al. [7] An R.S. model based on the ABOM approach is presented in this study. Initially, these authors demonstrated how user ratings might be derived using a deep learning technique. To further improve the recommendation system, the recovered features were used to generate aspect-based scores, which were then put into a tensor factorization engine. ABOM and rating prediction make up the majority of the proposed method. It utilised idle dirichlet allotment (LDA) to join the removed perspectives with the fundamental assessment terms to shape dormant viewpoints. Next, for each feature of the review, a score was calculated based on a linguistic technique. To further include the user's perspective on many elements and overall ratings, a three-dimensional T.F. approach is used to determine the representation of the underlying components. Hasan, K. M. A.et al. [9] these authors gathered information from reliable sources. The data were preprocessed to remove noise and improve usability. Kumar, P.et al. [12] indicated that a comprehensive examination of any issue might be undertaken by gathering a representative sample of Twitter user thoughts. Such an investigation might give helpful information to organizations or creators of movies, television shows, and businesses and warn them of an alarming change in the public's impression of their area. Identifying unfavourable tendencies early on may enable businesses to make more informed judgments about enhancing consumer satisfaction by concentrating on specific product characteristics. This study shows that the machine learning classifier utilized significantly impacts the inquiry's overall accuracy.

MATERIALS AND METHODS

Figure 1 is a flowchart that depicts the entire suggested methodology for processing a dataset and carrying out opinion categorization in Chapter III, which is concerned with the proposed approach. The flowchart has five primary stages, each of which is represented by a rectangular box with rounded sides and a number of connecting arrows. The first rectangular box on the flowchart's left side denotes the first stage, which entails preparing the dataset using Standard Scalar with Label Encoding. The second rectangular box represents the second stage, which entails utilising Improved LDA to tag parts of speech (PoS). The preprocessed dataset is trained using the Faster R-CNN with Bi-LSTM method in the third stage, which is depicted by the third rectangular box. Selecting is done at the fourth stage, which is symbolised by the fourth rectangular box.





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NORMALIZATIONUSING STANDARD SCALAR

Since their performance suffers if the underlying features do not closely match standard normally distributed data, some machine learning estimators demand that a dataset be standardised. Data is cleaned and then normalised for model testing and training. When data is divided, one set is used to test an algorithm, while the other is used to train it. The values found in the training data, together with the underlying logic and algorithms of the features, are used to build the training model. Consistency in all aspects is the aim of normalisation. Reduce the feature set's inherent variability by removing outliers and scaling everything down to a single unit. Here is how the average score is calculated:

$$Ds = z = \frac{x - \mu}{\sigma} - \dots$$
 (1)

POS TAG APPLYING USING IMPROVED LDA

Vectors for discrimination must be chosen. Not all discriminating vectors are created equally for pattern categorization. It is best to choose the vectors with the greatest Fisher discrimination values since they provide more details regarding dispersion both between and within classes. A desirable classification property that discrimination vectors must possess is statistical uncorrelation. Next, present a comprehensive set of recommendations utilising an enhanced LDA (ILDA) methodology. The remainder of the paper is divided into the following sections. This second section examines the three LDA advancements' theoretical bases. Normally, picture data is kept in the form of a two-dimensional (2-D) matrix (A*B), which may be transformed into a vector of size H using the equation H=A*B. As a consequence, this study could be able to draw out of the image library some typical dimensions.

FEATURE SELECTION

It is an estimator that fits randomised decision trees to distinct subsamples of a dataset to boost accuracy and control over data fitting. With 75 trees used in its design and 50 trees used in its construction, the Extra Trees Classifier had an accuracy rate for the test data of 90.27 percent and 90.73 percent, respectively. This model is not further developed since adding more trees does not significantly increase accuracy.

$$\varphi_i^* = \underset{\varphi i \in \rho}{\operatorname{argmin} \left\{ \sum_{j=1}^m l(yj, yj) \right\}}$$
(6)

$$= argmin\{\sum_{i=1}^{m} l(flightGBM(\varphi_i, x_j), y_j\} -----$$
 (7)

In Eq. (7), ℓ (\bullet) is the loss function, y is the prediction y j=flight GBM (ψ i, xj), and yj is the given actual class. Let k be a scalar of n bits, P and Q two points of an elliptically curve defined on a finite field P by the equation (:

$$E: Y^2 + a1XY + = X^3 + a_2X^2 + a_{4x}X + a_6 - - - -$$
 (8)

where a1; a2; a3; a4 2 F. In this work, to consider the finite prime fields Fp. Eq. (9) then becomes:

$$E: Y^2 = X^3 + aX + b - - - -$$
 (9)

OPINION CLASSIFICATION USING STACKING ENSEMBLE METHOD

Due to their superior performance versus individual learning models in a variety of tasks, including classification and regression, ensemble models have recently acquired popularity. Early 1990s ensemble learning research made the case that it was possible to combine a number of relatively weak learning algorithms to create more powerful ones. Using a single data set, many learner modules, and ensemble learning, various predictions are produced. A more accurate picture is provided by the sum of all expert predictions. The technique normally involves two steps. From the training data, a subset of students is chosen and included in a single prediction model. A number of forecasts were created utilising data that was distinct from that used in the underlying models to create the composite model.





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RESULTS AND DISCUSSION

The proposed method has implemented by using python programming language with spyder 3.8 framework. Figure 2 is a graph that illustrates reviews and analysis instances, illustrating both positive and negative types of input. The audits are displayed on the x-pivot, while the thickness of those surveys is displayed on the y-hub. The graph resembles a histogram, with bars of varying heights standing in for the number of reviews falling into various densities. The testimonials in Figure 2 probably refer to written evaluations or remarks made by consumers or users of a certain item or service. The density metric might be a score or rating determined using natural language processing (NLP) methods, which would be a measure of the sentiment or tone of those evaluations. The examination of punctuation for words is shown in figure 3. The graphic highlights both positive and negative evaluations of the product. This graph shows reviews along the x-pivot, and thickness is shown along the y-hub. The classification performance matrices for a variety of models, including DT, SVM, NB, Logistic Regression, Stacking Ensemble, are displayed in Table . Metrics including accuracy, precision, recall, and F-measure are included in the matrix. Stacking Ensemble model exceeds the other models in terms of accuracy, scoring 0.972 as opposed to 0.85 to 0.89 for the other models. The SVM and LR models had the greatest accuracy ratings (0.90), closely followed by the NB model (0.88). The Decision Tree and Stacking Ensemble models are closely behind the SVM and LR models in terms of recall, scoring 0.87 and 0.870, respectively.

CONCLUSION

The findings show that the proposed approach has a higher F1-score and is more accurate than alternatives. The suggested approach has a few benefits over the conventional approach. Second, by fine-tuning the text's immediate context, general properties, and relationships, it can manage complicated textual data. Last but not least, this approach may be applied in a variety of situations where sentiment analysis is required, such as marketing, customer service, and product development. Recommended to present a state of the art technique for assessment mining that mixes profound learning with customary AI strategies. To increase classification accuracy, the LSTM and CNN models extract text features and evaluate sentiment. his proposed system was assessed on a few datasets and contrasted with the latest cutting edge philosophies.

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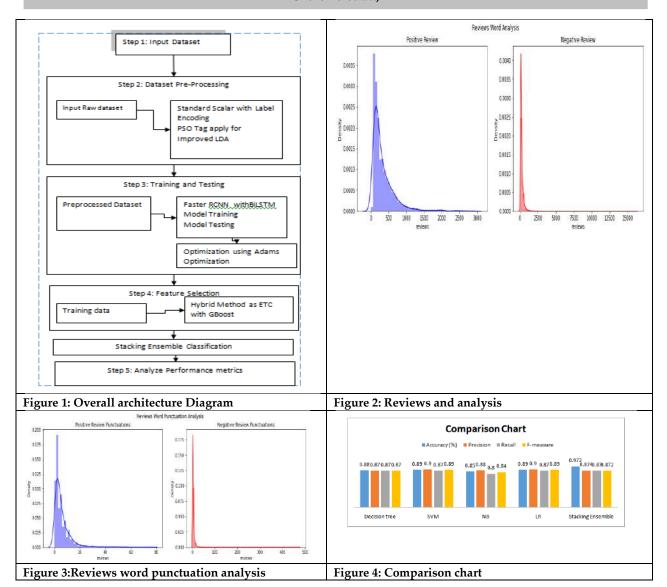
Table 1: Training and validation accuracy and loss

Epoch	Training Loss	Validation Loss	Training Accuracy	Testing Accuracy
1	0.1544	0.0679	0.9547	0.9779
2	0.0537	0.0526	0.9837	0.9820
3	0.0352	0.0596	0.9890	0.9826
4	0.0243	0.0465	0.9922	0.9848





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RESEARCH ARTICLE

An AI Based Keystroke Dynamics for User Authentication

Logidha V1* and Karthik S2

¹UG Student, Department of Information Technology, Sri Ramakrishna College of Arts & Science, Coimbatore, Tamil Nadu, India.

²Assistant Professor, Department of Information Technology, Sri Ramakrishna College of Arts & Science, Coimbatore, Tamil Nadu, India.

Received: 22 Feb 2023 Revised: 15 Mar 2024 Accepted: 21 Mar 2024

*Address for Correspondence

Karthik S

Assistant Professor, Department of Information Technology, Sri Ramakrishna College of Arts & Science, Coimbatore, Tamil Nadu, India.

Email: karthik@srcas.ac.in



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ABSTRACT

Authentication is frequently referred as the most critical part of a computer system security. Users commonly identify themselves using a combination of username and password, but sometimes this is not enough. Concerning web-based services, attacks like phishing or social engineering can easily result in identity theft. In addition, the widespread use of single sign-on services can seriously increase the consequences of such attacks. In these circumstances strong authentication is mandatory. Strong authentication is often implemented using additional authentication steps or specialized hardware modules, which is not suitable for web-based systems. However, biometrics can used to overcome these limitations. More specifically, behavioural biometrics based on keyboard typing patterns can provide an extra security layer on top of conventional authentication methods, with no additional cost and no impact to the user experience. This work aims to evaluate the feasibility of the implementation of strong authentication on the web using keystroke dynamics. This is carried out through the creation of a application prototype using python environment.

Keywords: Keystroke Dynamics, Behavioural Biometrics, Authentication, Keyboard Typing, Security.

INTRODUCTION

Our dependence on computers and digital platforms has been observed to be overwhelmingly increased to simplify our lives. The use of such automated information systems together has resulted in improved performance of the available networking services in the form of reliability and computational costs. With such efficient utilization the





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advances in technology have generated a collective interest in global access to such online platforms. However, at the same time there is a rise in the threats regarding to the security of computers. Usage of advanced methodologies to safeguard the system from attacks and frauds come under the topmost concerning priority of many research scientists. Hence there is a need to generate foolproof measures to prevent such unauthorized access is being worked upon. One such preventive method to give access to individuals by detecting their unique and behavioral pattern is an individual's typing rhythm. This unique typing rhythm tends to become a natural choice for security of computers and is commonly known as Keystroke Dynamics. It is observed that when a person types; the placement of his fingers, applied pressure on the keys and the regularly typed strings appears to be consistent for a specific individual. Hence this concept is used to differentiate between an intruder and a legit user as they will be typing on the keyboard anyway. Therefore, making such kind of typing rhythms easily accessible to track computer activities.

Background

The initial step to prevent any form of unauthorized control into a system is user authentication. This is the only operation that confirms the identity of an individual. This identity is matched with a pre-generated code or a registration number with the legit individual and is used in the process of identification. Once this identification is accomplished some form of indicator is shared to give access control to the user. Knowledge-based encryption is based on the idea of two people sharing their secrets. Username and Login credentials are two well-known examples. Whereas object-based authorization is characterized based on possession of specific things and relies on the concept of something which someone has. On the other hand, biometric based authentication is characterized by behavioral features an individual would have. In real time, the classifications of object and knowledge based are merged to fulfill the process of authentications such bank passwords and their PINS. One major disadvantage with regards to this classification-based authentication is the ability to memorize and manage multiple such PINS and recalling them. Therefore, the usage of biometrics authentication is preferred as it overcomes these issues and makes use of automated methods to identify and verify the individual. Also, this form of authentication is gaining worldwide popularity as they provide an extra level of security.

Motivation

User authentication has always been a fundamental component of access control. Nowadays, the Internet is the standard platform for communication and a tremendous amount of sensitive data is transmitted between computer systems. This makes the process of identity verification more important than ever. However, basic username and password combination is used the same way we did a decade ago. Attacks like man-in-the-middle, phishing or social engineering can easily result in identity theft. This can lead to access to private information, exploitation of trust relationships and other criminal activities. Furthermore, with the widespread use of single sign-on services, where a single set of credentials can authenticate users in multiple websites, consequences of identity theft can be devastating. Even though in most cases simple password-based authentication will suffice, other environments like online commerce and banking platforms can definitely benefit from strong authentication to ensure proper protection of sensitive, private and confidential data. Strong authentication doesn't have a standard definition, but it generally aims to deliver increased security beyond standard authentication methods. It is often implemented with two-factor authentication and usually based in knowledge or token-based approaches. Although these methods can effectively improve security in authentication, what is really validated are passwords, tokens and keys. Also, user credentials can be stolen, lost, shared or manipulated, resulting in compromised security. Biometrics can be used to overcome the limitations imposed by classical solutions by allowing identity verification based on the user himself. Unfortunately, authentication solutions based on biometrics may not always be easily deployed, since most of them require specific hardware (e.g. fingerprint reader). Keystroke dynamics, a behavioural biometric, presents itself as a viable contender to implement strong authentication on the web. It has the distinct advantage of not requiring specific hardware, while maintaining the desirable properties of the more mainstream biometric solutions. Also, as we will explore in this work, it can be seamlessly integrated with existing password-based authentication.





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Problem Statement

One of the major issues or a problem the digital world face is with regards to passwords used. Although, a few organizations do exist that provide secured authentication for users when they login but on the other hand there are still most sites that skip the process of identification leading to major crimes across the globe. Hence the research work in this thesis aims to suggest better security systems that could be analysed using the typing behaviour of individuals using keystroke dynamics. Main emphasis is being put on to detect the typing behaviour of individuals using the concepts of validate the keystrokes timing features of users and give them a more secure and efficient system than ever before.

Objectives

This work aims to evaluate the viability of keystroke dynamics as a method for implementing strong authentication on a web-based environment. It should result in an implementation capable of enrolment and identity verification, applying typing behaviour on top of basic username and password credentials. The goal of this study is to compare the performance of some classical matching algorithms proposed by previous research. Although numerous experiments have achieved encouraging results, there is significant variation in the reported effectiveness, which makes it hard to assess the full potential of keystroke dynamics as a biometric for authentication. We'll also look at how adaptive techniques can influence the performance of keystroke dynamics. Finally, most studies on keystroke dynamics tend to use a single passphrase for all users; this work's experiment should account for individual authentication credentials for every user, which is arguably a better representation of the real world.

SYSTEM STUDY

System analysis will be performed to determine if it is flexible to design information based on policies and plans of organization and on user requirements and to eliminate the weakness of present system. This chapter discusses the existing system, proposed system and highlights of the system requirements.

EXISTING SYSTEM

Distortion in keystroke dynamics is not a well-researched field, and we did not find many results when doing literature search. However, it measured two typing samples of keystroke dynamics data and used two different measures to compare the samples. However, one method that could be used for detecting distorted timing information is Benford's Law and ZIPF's Law. Benford's Law, or the first-digit law, is an observation in a set of numerical data where the first digit, or leading digit, is more likely to be small. In a balanced distribution of numbers between 1 and 9 there would be exactly 11% for each number to be the leading digit. However, if Benford's Law is obeyed then the change of the leading bits to be small increases.

DISADVANTAGES OF EXISTING SYSTEM

However, the results showed that only latency values from keystroke dynamics timing information followed the law. While duration values did not follow the law. More resources used for re-keying because it is done for each join or leave operations.

PROPOSED SYSTEM

In order to authenticate a user using keystroke dynamics we need to create a reference average value for each user that will represent, as accurately as possible, their specific typing behaviour. This template varies a lot depending on whether static or continuous authentication is used. In authentication we want to create a template that reflects the typing rhythm that the genuine user uses in order to type the password. This average value is created based on enrolment samples, where the user would get requested to type their password a number of times. The features, such as duration and latency, are then extracted and the average typing rhythm is calculated and stored as a reference average. When a user tries to authenticate, the system will check their typing rhythm, which is referred to as a probe, against the reference template and then either reject or accept the user based on a criterion. This criteria





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for decision making are decided by a threshold which is created by a minimum and maximum average time in milliseconds.

ADVANTAGES

The proposed Keystroke dynamics is based on the assumption that each user can be authenticated because of their unique typing manner. This is because keystroke dynamics performs on a millisecond's precision level meaning it is impossible to accurately recreate the way another user types. This is true even for a user who is typing their own password, as they would not be able to type exactly the same way they did last time. Even though the user might type one of the keys or key pairs the same, there are still other keys that they could type in a different way. It is because of this reason keystroke dynamics works as an authentication method.

SYSTEM ANALYSIS AND DESIGN

MODULE DESIGN

Keystroke timing

From the Key Down and Key Up time of each keystroke we can calculate the duration and latency of a key. The duration of a key is how long the key was held down, this can also be referred to as dwell or hold time. While the latency of a key is the time between releasing one key and pressing another key, and this can sometimes be referred to as flight time. We differentiate between 4 different latencies, given as:

- 1. •pp-latency: The timing it takes to press down one key and the next key.
- 2. •rr-latency: The timing it takes to release one key and the next key.
- 3. •rp-latency: The timing it takes to release one key and press the next key.
- 4. •pr-latency: The timing it takes to press down one key and release the next key.

In order to get the pp-, rr- and pr-latency we have to use the timing information from duration and rp-latency. We can calculate pp-latency as *latpp= durA+latrp*, rr-latency as *latr r = latrp+ durB* and pr-latency as *latpr= durA+ latrp+ durB* where *durA* and *durB* represents the duration of two different keys. The following figure shows the timing values we can extract if a user types the keys *A* and *B*. From these latencies, only the rr-latency and rp-latency can be negative. For example, for the rr-latency we can press the shift key, followed by pressing the C key, and then release the C key before releasing the shift key. The same can be said with rp-latency as we can press the next key before releasing the previous key, for example we can press the C key before releasing the shift key. By using pp-latency, rp-latency and duration for our timing values. The naming of pp-latency refers to press-press-latency, however, we will refer to this as Key Down-Key Down latency (DD). While rplatency refers to press-release-latency, which will be called KeyUp-Key Down latency (UD).

Matching module

In this module, extracted timing features are compared against those stored in the database, using a matching algorithm from statisticalt approaches. Ultimately, this process results in a matching score, that represents a similarity measure between the extracted features and the ones previously stored in the database.

Decision module

Given the matching score and a decision threshold, this module is responsible for either accepting or rejecting the claimed user identity, based on the matching score and a predefined threshold.

CONCLUSION AND FUTURE ENHANCEMENT

With the conclusion of this work, and regarding the original research goals, keystroke dynamics can be considered a viable choice to implement strong authentication on the web. We've shown that this technology can be integrated on top of a tradition authentication procedure, taking advantage of user input to seamlessly capture and classify the





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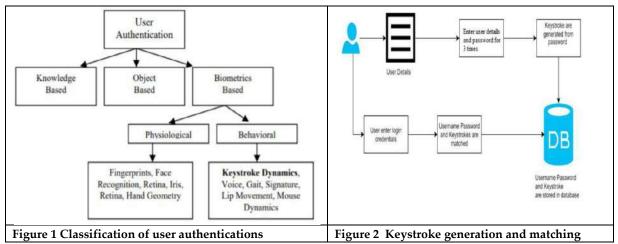
associated typing behaviour. Keystroke dynamics presented some notable advantages over other alternatives. No specialized hardware was needed. Biometric data acquisition was completely transparent and multifactor authentication was effectively achieved with a single conscious action from the user. In addition, even simple matching algorithms, as the ones used in this study, can yield reasonable accuracy on authentication. Although the proposed goals of this work were successfully met, some points could have been improved. An assessment of the system usability should have been done by the users that participated in the study. Furthermore, the target population was greatly biased, mainly computer science students, which are probably better typists than the average individual. Although more complex, applying this study to a more diverse population would have been a closer representation of a real environment.

SCOPE FOR FUTURE ENHANCEMENT

For instance, it would be interesting to develop a complementary study, focused on the evaluation of more sophisticated algorithms, in order to further improve matching accuracy.

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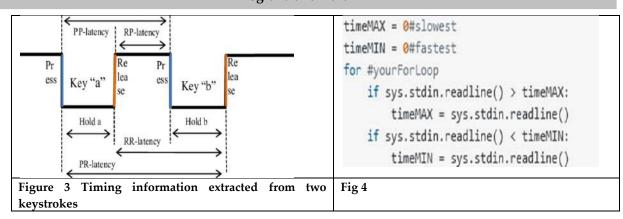
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RESEARCH ARTICLE

IoT Awarecensored Regressive Gaussian Kernelized Recurrent Deep Neural **Network for Automatic Agriculture Prediction**

R.Hemalatha¹ and G.Nivashini^{2*}

¹Research Supervisor, PG & Research Department of Computer Science, Tiruppur Kumaran College For Women, Tiruppur, Tamil Nadu, India.

²Research Scholar (Ph.D) PG & Research Department of Computer Science, Tiruppur Kumaran College For Women, Tiruppur, Tamil Nadu, India.

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*Address for Correspondence

G.Nivashini

Research Scholar (Ph.D) PG & Research, Department of Computer Science, Tiruppur Kumaran College For Women, Tiruppur, Tamil Nadu, India Email: nivashini96phd@gmail.com



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ABSTRACT

Agriculture plays a vital role in the country's economy. The most widespread concern among farmers is that they do not choose their crops based on the requirements of the soil conditions. The majority of farmers are unaware of the significance of planting crops at the appropriate time and place. Due to these agricultural practices, seasonal climatic conditions are altering, resulting in food insecurity. Therefore, agricultural prediction helps farmers make decisions about the farming approach to increase crop yield. Various deep learning or machine learning techniques have been developed to increase the quantity and quality of production from crop fields. However, achieving accurate and time-efficient predictions is a significant concern. In this paper, a novel method named Censored Regressive Gaussian Kernelized Gradient Recurrent Neural Network (CRGKGRNN) is developed for accurately predicting crop planting in the agriculture domain, aiming to enhance crop productivity. The CRGKGRNN method comprises four distinct processes such as data acquisition, pre-processing, feature selection, and classification. In the data acquisition phase, IoT sensors are deployed in the agricultural land to collect soil moisture, temperature, humidity, potassium, phosphorus, nitrogen, and rainfall. The data collected by IoT sensors are pre-processed in the second step to address missing values or null, as well as noise. Damped Least-Squares (DLS) method is utilized to identify and remove duplicate or noisy data. In the third step, attribute selection is performed using Sokal-Michener's simple matching indexive censored regression to choose a subset of relevant features and remove irrelevant or redundant attributes, thereby minimizing the time consumption of predictions. Finally, the Gaussian Kernelized Adaptive Gradient Recurrent Neural Network is applied for prediction process which comprises three layers namely the input layer, two or more hidden layers, and the output layer.





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Keywords: Agriculture, IoT, Crop prediction, Damped Least-Squares (DLS), Recurrent Neural Network, Prediction

INTRODUCTION

Agriculture is the fundamental practice of cultivating soil, growing crops, and increasing productivity. It is deeply influenced by climate variables such as temperature, precipitation, humidity, and sunlight. Changes in these factors directly affect crop growth, yield, and overall agricultural productivity. Recognizing the intricate relationship between climate conditions and crop suitability, advanced technologies are required for predictive modelling to provide valuable insights for decision-making in farming practices. Although numerous machine learning models have been developed for identifying crops but their performance has not been effectively investigated. The Interfused Machine Learning with Advanced Stacking Ensemble model (IML-ASE) was developed in [1], aims to improve crop prediction accuracy by leveraging information related to farming, climate, and soil conditions. This enhancement enables farmers to make informed decisions regarding crop selection for cultivation. However, it noted that the designed method did not succeed in minimizing the processing time. A deep reinforcement learning (DRL)based crop classification system was developed [2] to address the challenges faced by cultivation in precision agriculture. However, the performance of accurate crop classification was not performed. A smart agriculture system based on artificial intelligence and deep learning was developed in [3]. However, addressing the time complexity associated with smart agriculture proved to be a challenging task. The Agriculture Factor-based Relevance Vector Analysis Model was developed in [4] to predict optimal crops and enhance productivity. However, a significant challenge occurred due to the high error rate in crop prediction. Various machine learning algorithms were developed in [5] for precise crop prediction. But, it was not efficient for evaluating large crop data from diverse geographic regions using IoT. Feature selection and classification techniques were developed in [6] to enhance the prediction accuracy of plant cultivations. However, the issue of prediction time performance remained unresolved. A crop recommendation system, developed was in [7], utilizes Map Reduce and K-means clustering. However, it failed to deliver more accurate recommendations. A Fuzzy Enumeration Crop Prediction Algorithm (FECPA) was developed in [8] to enhance the accuracy of predicting suitable crops and improve productivity. However, it encountered limitations in achieving accurate predictions with minimal time due to the absence of machine learning algorithms. Recurrent Neural Networks (RNNs) were developed in [9] to determine the most probable scenarios for multi-temporal crop prediction.

RELATED WORKS

A graph neural network model was introduced in [11] for crop suitability evaluation with higher accuracy. However, it did not succeed in significantly reducing the cost of trial and error in the crop suitability evaluation process. A quantum value-based gravitational search algorithm (GSA) was developed in [12] to predict the suitable soil for optimal crop production. However, the implemented algorithm proved ineffective when applied to real-time optimal crop selection systems. Microclimate modelling techniques were developed in [13] with the aim of predicting future crop suitability. However, attaining higher accuracy posed a significant challenge in forecasting future crop suitability. A transfer learning approach was introduced in [14] for agricultural crop classification. However, this method did not incorporating a more sophisticated approach to enhance overall accuracy. A statistical and machine learning approach was introduced in [15] for predicting crop suitability types. However, the crop selection methodology did not use any deep learning techniques for improving accuracy. An LSTM RNN model was developed in for crop selection based on weather conditions and soil parameters. However, it did not integrate sensors to gather real-time, more precise data on weather conditions and soil parameters for improving the model's efficiency.





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PROPOSED METHODOLOGY

Crop suitability prediction in smart agriculture is used to assist farmers in predicting specific seasonal crops according to the weather conditions. As weather conditions continue to change, crop prediction becomes more essential for improving crop productivity. Based on this motivation, a novel CRGKGRNN is introduced in this section for accurate crop prediction with minimum error as well as minimal time consumption. Figure 1 illustrates the architecture of the proposed CRGKGRNN method, comprising four major processes: Data acquisition, preprocessing, feature selection, and classification, designed to enhance the accuracy of crop prediction. Initially, IoT sensors are deployed in agricultural fields to collect soil characteristics, including, temperature, humidity, potassium, phosphorus, nitrogen, and rainfall. These data are stored in a dataset with features denoted as $A_1, A_2, A_3, ... A_m$ and training instances represented as $SD_1, SD_2, SD_3, ... SD_n$. The first step involves data pre-processing to clean and transform raw data, addressing null values and removing duplicate values. Subsequently, the feature selection process is performed using Sokal–Michener's simple matching index and censored regression to identify significant features for efficient prediction within minimal time. Finally, the proposed CRGKGRNN method utilizes Gaussian Kernelized Adaptive Gradient Recurrent Neural Network to enhance crop prediction performance, achieving higher accuracy and minimizing errors. The distinct processes of the proposed CRGKGRNN are described in the following subsections.

Data acquisition phase

The data acquisition phase marks the commencement of the proposed CRGKGRNN, involving the collection of data from the Smart Farming using Machine Learning dataset. This dataset encompasses information on soil and environmental characteristics, including nitrogen, phosphorus, potassium, temperature, humidity, rainfall, etc. The data is presented in both numeric and categorical formats. The dataset's structure is defined by rows and columns, providing a comprehensive representation of the information.

Data preprocessing

The second step of the proposed technique is the data pre-processing that involves cleaning and transforming raw data into a suitable format for analysis. This phase is important for improving the quality of the data, handling missing values, and removes the duplicate data making it compatible with machine learning algorithms.

Proportional Weighted Average Sampling based null values handling

Null values, also known as missing values, are data points that have not been recorded or are unavailable in a particular cell of the dataset. Dealing with null values is a fundamental aspect of data pre-processing. The proposed CRGKGRNN utilizes the Proportional Weighted Average Sampling technique in which null samples from a larger population are chosen. To start with the raw input dataset 'DB' and formulated in the form of matrix as given below.

$$I = \begin{bmatrix} A_1 & A_2 & \dots & A_m \\ SD_{11} & SD_{12} & \dots & SD_{1n} \\ SD_{21} & SD_{22} & \dots & SD_{2n} \\ \vdots & \vdots & \dots & \vdots \\ SD_{m1} & SD_{m2} & \dots & SD_{mn} \end{bmatrix}$$
(1)

Where, 'm' column features $A_1, A_2, A_3, ... A_m$ are present with overall sample instances of 'n row $SD_1, SD_2, SD_3, ... SD_n$ respectively. The Proportional Weighted Average Sampling method determines a null value by taking a weighted average of a certain number of consecutive observed values in the particular column. The formula for calculating the Proportional Weighted Average is given below,

$$P_{WA} = \frac{\sum_{i=1}^{n} SD_{i} * \beta_{i}}{\sum_{i=1}^{n} \beta_{i}}$$
(2)

Where, P_{WA} denotes a proportional Weighted Average for the missing value, SD_i denotes an observed value for data point, β_i denotes a weight assigned to data point SD_i . Therefore, the formula provides a way to estimate missing values based on the available data points in the dataset.





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// Algorithm 1: Data pre-processing

Input: Dataset 'DB', features $A = \{A_1, A_2, A_3, ... A_m\}$ and data samples $SD_1, SD_2, SD_3, ... SD_n$

Output: Pre-processed dataset

Begin

- 1. For each Dataset 'DB' with Features 'A'
- 2. Formulate input vector matrix I' as given in (1)
- 3. **If** missing feature value **then**
- 4. Measure proportional weighted average between the feature as given in (2)
- 5. Fill the feature value to the respective column
- 6. End if
- 7. **for each** data point
- 8. Measure the squared differences between the values using (4)
- 9: **if** $\arg \min \sum_{i=1}^{n} [f_i(x)]^2$ **then**
- 10. Normal data
- 11: **else**
- 12: Duplicate data
- 13. End if
- 14. Remove Duplicate data
- 15. End for

End

Algorithm 1 outlines a step-by-step process of data pre-processing. Initially, the number of features and raw data is gathered from the dataset. Following data collection, any missing or null values are addressed using a proportional weighted average sampling method. The null values are replaced with the proportional weighted average value. Subsequently, the DLS method is utilized to analyze the data points in a specific column. If the deviation is minimal, the data point is considered normal. On the other hand, if the deviation is large, the data point is identified as a duplicate. Finally, a structured dataset is obtained.

// Algorithm 2: Sokal-Michener's simple matching indexive censored regression

Input: pre-processed Dataset 'DB', features $A = \{A_1, A_2, A_3, ... A_m\}$ and data samples $SD_1, SD_2, SD_3, ... SD_n$

Output: relevant feature selection

Begin

1. For each Dataset 'DB' with Features 'A'

2Measure Sokal-Michener's simple matching using (5)

- 4. Formulate censored regression outcome to obtain relevant features as given in (6)
- 5. If (SM > T) then
- 6. **Select**relevant features
- 7. else if (SM < T)then
- 9. **Remove ir**relevant features
- 10. **End if**
- 13. End for
- 14. Return (relevant features)

End

Algorithm 2, illustrates the process of selecting significant features to minimize the time complexity of crop prediction. The pre-processed dataset serves as input for Sokal–Michener's simple matching indexive censored regression. Initially, the algorithm assesses the relationship between features by employing Sokal–Michener's simple matching index. Subsequently, censored regression is applied for measuring the relationship, aiming to identify relevant features by setting a predetermined threshold. Finally, the outcomes of the regression provide the highly relevant features, which are then chosen for crop prediction. Figure 1 illustrates the architecture of the proposed





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CRGKGRNN method, comprising four major processes: Data acquisition, pre-processing, feature selection, and classification, designed to enhance the accuracy of crop prediction. Initially, IoT sensors are deployed in agricultural fields to collect soil characteristics, including, temperature, humidity, potassium, phosphorus, nitrogen, and rainfall. These data are stored in a dataset with features denoted as $A_1, A_2, A_3, ... A_m$ and training instances represented as $SD_1, SD_2, SD_3, ... SD_n$.

The hidden layer captures information from previous layer where Gaussian kernel process is applied for analysing the testing and training data as given below,

$$GK = \sum_{i=1}^{n} \sum_{j=1}^{m} \exp\left[-0.5 * \frac{(|SD_{t} - SD_{r}|^{2})}{v^{2}}\right]$$
 (7)

Where, GK indicates a Gaussian kernel between the testing data point DS_t and the training data point ' DS_r ' and 'v' indicates a deviation. Based on the analysis, the data points are classified into different classes. The output of the hidden or recurrent layer is formulated as follows,

$$R_t = F\left(\alpha_{ih} \cdot SD_t + \alpha_{hh} \cdot R_{t-1} + B_R\right) \tag{8}$$

Where, R_t denotes a hidden or recurrent layer output at time't', F indicates sigmoid activation function, α_{ih} denotes a weight matrices for the input and hidden state, SD_t indicates a input data sample at time step't', α_{hh} indicates a weight matrices for the hidden state, R_{t-1} indicates a previous hidden state at time step 't-1', B_R indicates a bias term in hidden layer that stores the value '1'. Once all the time steps are completed the final current state is used to calculate the output. The output is generated as follows,

$$Y_k(t) = F(\alpha_{ho}.R_t + B_o) \tag{9}$$

Where, $Y_k(t)$ denotes actual classification outcome at time 't', F indicates sigmoid activation function, α_{ho} denotes a weight matrices from the hidden state to output state, R_t indicates a hidden layer output, B_o indicates a bias term in output layer that stores the value '1'. After that, error is computed based on the expected and actual classification outcome.

$$e = [Y_E(t) - Y_k(t)] \tag{10}$$

Above depicts a graphical illustration of the accuracy of crop prediction using three different methods CRGKGRNN and existing IML-ASE [1], DRL [2], with respect to number of data samples ranging from 200 to 2000. In Figure 3, the x-axis denotes the number of data samples, and the y-axis indicates the crop prediction accuracy. The graphical effects show that the proposed CRGKGRNN achieved higher crop prediction accuracy compared to existing methods [1] and [2], respectively.

CONCLUSION

Smart farming is a modern approach that utilizes digital and IoT devices in conjunction with machine learning to enhance productivity and optimize resource utilization. The IoT-enabled crop prediction system offers numerous advantages to farmers, enabling them to make informed decisions for cultivation, optimize resource usage, and implement precision farming techniques. In this paper, we introduce an efficient crop prediction method called CRGKGRNN, designed to predict suitable crops for cultivation with higher accuracy and minimal time consumption. The main objective of CRGKGRNN is to improve the accuracy of crop prediction. Following data preprocessing, relevant features are selected, and a classification of different types of crops is presented to achieve accurate predictions with minimal error. The experimental evaluation of CRGKGRNN highlights its superior performance. Specifically, it increased accuracy in crop prediction, precision, recall, and F1-score, while simultaneously reducing the time required for crop prediction compared to existing methods.





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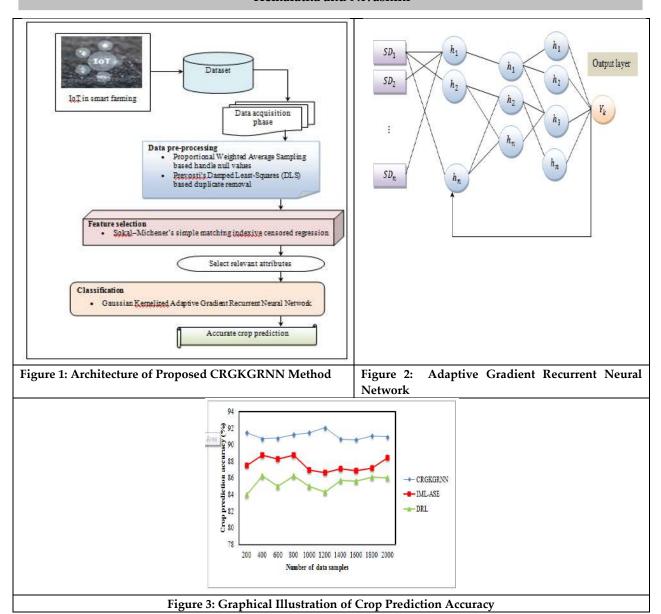
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RESEARCH ARTICLE

AI Based Ayurvedic Plant Divination

Usha M S *1, Navyashree S2 and Pratheek Raj Urs C P2

¹Associate Professor, Department of Computer Science and Engineering, NIE North, Mysuru-Karnataka-India.

²Department of Computer Science and Engineering, NIEIT, Mysuru, Karnataka, India.

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*Address for Correspondence

Usha MS

Associate Professor,

Department of Computer Science and Engineering,

NIE North, Mysuru-Karnataka-India.

Email: ushams@nie.ac.in



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ABSTRACT

In the convoluted realm of medicinal plant utilization, the fusion of ancient wisdom and modern technology emerges as an imperative intervention. Celebrated for its efficiency in treating chronic illnesses. Using Tensor Flow Lite and a Convolutional Neural Network (CNN) algorithm, the application revolutionizes Ayurvedic plant identification through advanced image processing. The user-friendly app, developed in Android Studio, facilitates rapid, precise scanning and detection, providing detailed insights into medicinal benefits, traditional uses, and cultural significance. Beyond recognition, the application encourages active user engagement in cultivation practices and conservation efforts. This innovative tool is essential to safeguard endangered Ayurveda plants, ensuring the preservation of their rich heritage. The project offers a synergistic solution, merging ancient wisdom with cutting-edge technology to conserve Ayurvedic botanical treasures for future generations. This contributes to the intersection of traditional knowledge and technological solutions in a concise and impactful manner, addressing the critical need for the preservation of Ayurvedic plant species.

Keywords: Medicinal plants, Ayurveda, Computer vision, Traditional medicine, Medicinal properties

INTRODUCTION

In the realm of healing traditions, Ayurveda, an ancient medicinal system originating in India, has provided profound solutions for diverse ailments by harnessing the potent properties inherent in medicinal plants. Rooted in a holistic approach that considers the interconnectedness of mind, body, and spirit, Ayurveda boasts a history spanning thousands of years. Despite their historical significance, awareness of Ayurvedic botanical remedies has waned in contemporary society, necessitating a concerted effort for preservation and reintroduction. This research





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endeavors to address this imperative through the development of an innovative system at the confluence of ancient wisdom and state-of-the-art technology. Leveraging a Convolutional Neural Network (CNN) algorithm implemented via TensorFlow and seamlessly integrated into a Mobile Application developed using Android Studio. The project offers a comprehensive guide to Ayurvedic plants. This application not only visually identifies the plants but also provides extensive information regarding their benefits and methods of incorporation into daily life. Central to this system is a sophisticated computer vision mechanism empowered by advanced algorithms. This mechanism captures and processes intricate leaf images, extracting features such as eccentricity, color and shape. The automated plant identification feature, akin to a digital herbalist, functions as a scanner, enabling users to effortlessly detect and learn about various Ayurvedic plants. This fusion of Artificial Intelligence and Ayurvedic knowledge not only bridges awareness gaps but also nurtures a deeper connection between individuals and the healing properties of nature. Additionally, this project draws inspiration from recent advancements in automated plant identification. From algorithms utilizing Gaussian distribution for precise classification to innovative combinations of Speeded Up Robust Features (SURF) and Histogram of Oriented Gradients (HOG) achieving near-perfect accuracy rates, these technological strides underscore the potential for a cultural resurgence. This convergence represents not only a technological leap but also a harmonious integration where ancient wisdom synergizes with modern scientific methodologies. By seamlessly blending tradition with innovation, this transformative system revitalizes the understanding of Ayurvedic plants, ensuring the enduring legacy of their healing properties. This endeavor transcends technological innovation; it signifies a cultural revival where the echoes of Ayurveda resonate through the corridors of modernity, emphasizing the invaluable treasures nature provides for holistic well-being. This research contributes to the ongoing discourse in the intersection of traditional medicinal knowledge and contemporary technological solutions, offering a promising avenue for the preservation and dissemination of Ayurvedic wisdom.

LITERATURE REVIEW

The intersection of machine learning, deep learning and Image processing has seen significant exploration in the domain of medicinal plant classification, particularly employing convolutional neural networks (CNNs). Previous research has predominantly relied on CNNs for the recognition of plant species based on leaf characteristics, demonstrating commendable accuracy in various studies. However, inherent limitations, particularly in handling complex backgrounds and smaller leaves observed in the papers proposed by R. Upendar Rao et al., have prompted the need for advancements in this field. Our proposed application represents a substantial leap beyond these constraints, incorporating state-of-the-art technologies such as Tensor Flow and advanced CNN architectures. A critical analysis of prior literature projects proposed by Nilesh Bhelkar et al., that the algorithms employed were often deemed insufficient and inaccurate, impeding their overall efficiency. In stark contrast, our application employs CNN, achieving an outstanding accuracy of 94%, thereby transcending the limitations of prior projects proposed by Dinesh Shitole et al., and Rakibul Sk et al., significantly enhancing efficiency in Ayurvedic plant recognition. The distinctiveness of the application lies in its user-centric features, designed to facilitate active engagement in plant identification and learning. Noteworthy is the incorporation of a comprehensive database, setting our application apart by providing users with extensive information beyond mere plant identification. This synthesis of advanced technology precision and user-centric design collectively positions our application as a pioneering and distinctive solution in the landscape of Ayurvedic plant recognition. In conclusion, while earlier projects proposed by Fadil Chady et al., laid the foundational groundwork, proposed application signifies a substantial advancement, overcoming the inefficiencies of past algorithms and delivering a more accurate and efficient platform for Ayurvedic plant classification. This research contribution establishes a new paradigm, fostering a seamless blend of technological sophistication and user engagement in the exploration and dissemination of herbal medicine knowledge.





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Problem Statement

In the realm of natural remedies, Ayurvedic plants pose a significant challenge for recognition and accessibility, attributed to complex identification methods and the inadequacies of existing plant recognition apps. The intricate nature of Ayurvedic knowledge exacerbates this issue, leading to a substantial gap in understanding and utilization. Our research identifies this pressing problem and endeavors to bridge the gap by employing artificial intelligence. Through the development of a user-friendly application, we aim to empower individuals spanning from enthusiasts to practitioners, to overcome the hurdles of Ayurvedic plant recognition. The objective is to create a comprehensive solution that not only addresses the complexities of identification but also fosters a deeper connection with the healing potential of Ayurvedic flora.

PROPOSED METHODOLOGY

This innovative application's methodology is centered around cutting-edge technologies, prominently computer vision and deep learning algorithms, facilitating the scanning, identification, and comprehensive analysis of Ayurvedic plants. The process begins with the capture of high-resolution plant images, subjected to pre-processing techniques to enhance clarity and quality. Leveraging the Convolutional Neural Networks (CNN), a robust deep learning algorithm, the application conducts intricate plant recognition, discerning unique features such as leaf patterns, shapes, and textures. Implemented in Python using TensorFlow, the CNN algorithm plays a central role in the precise identification of Ayurvedic plants. TensorFlow provides a robust framework for the development and training of deep neural networks, enhancing the efficiency and accuracy of the identification process. The integration of Android Studio, utilizing Kotlin, facilitates seamless application development for Android devices. Upon successful identification specialized algorithms are deployed for image detection and processing employing complex pattern recognition methodologies to extract essential characteristics from the plant images. The CNN architecture that is designed for image classification tasks, ensures precise feature extraction contributing to accurate plant classification. Subsequent to identification, the application retrieves comprehensive information about the recognized plant from an extensive database.

This database includes the plant's local name, uses, different properties and botanical information. The intention is to empower common users with the ability to easily identify and understand Ayurvedic plants. This information is presented in distinct sections within the application, each dedicated to specific aspects such as plant recognition and detailed analysis of medicinal properties. The app operates by performing all algorithms in the backend, and upon completion, it displays the output by presenting the matching plant's information, including its uses. In cases where the scanned plant is not identified as an Ayurvedic plant, the application connects to Google and provides relevant results, indicating that it is not an Ayurvedaplant. The significance of utilizing CNN, TensorFlow and Android Studio lies in their combined efficiency. CNN excels in image recognition tasks, TensorFlow provides a robust deep learning framework, and Android Studio facilitates user-friendly application development. Python serves as a versatile language for seamless implementation. The meticulous design of the application ensures not only the accurate scanning and identification of Ayurvedic plants but also the dissemination of valuable information, fostering a deeper understanding of these natural resources in the context of holistic healthcare.

Architecture

In the proposed research application, users initiate plant identification by capturing an image, subsequently undergoing pre-processing for image enhancement. The Convolutional Neural Network (CNN) then scrutinizes the extracted features for precise plant recognition. Upon successful identification, comprehensive plant information is retrieved from a dedicated database. The user-friendly interface systematically categorizes and presents this information, enabling user exploration. Users engage further by interacting with the application, saving, and sharing plant profiles, fostering a deeper understanding of Ayurvedic plants. This architecture seamlessly integrates image processing, deep learning, and database retrieval, offering an efficient and user-friendly solution for Ayurvedic plant recognition.





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Work Flow

Image Capture: High-resolution images of Ayurvedic plants are captured.

Pre-processing: Images undergo enhancement and noise removal.

Feature Extraction: Key plant features like patterns and textures are extracted.

CNN Implementation: Convolutional Neural Network analyzes features for plant recognition. Plant Identification: CNN matches features with a trained dataset for accurate identification.

Information Retrieval: Comprehensive data, including medicinal benefits, is retrieved from a database.

Data Presentation: User-friendly interface categorizes and presents plant information.

User Interaction: Users can explore, save, and share plant profiles.

RESULT ANALYSIS

Comparison Of Existing Models Vs Proposed Model

CONCLUSION

In conclusion, the work is poised to address identified shortcomings in Ayurvedic plant classification through an innovative application. Through a thorough literature survey, challenges in handling complex backgrounds and achieving precise plant recognition have been recognized, paving the way for the proposed solution. The forthcoming application will represent a groundbreaking leap forward by leveraging cutting-edge technologies, including TensorFlow Lite and advanced CNN architectures. The critical analysis undertaken in this research underscores a commitment to overcoming algorithmic inefficiencies and ensuring enhanced accuracy and efficiency in Ayurvedic plant recognition. The uniqueness of the future application will be evident in its user-centric features and the integration of a comprehensive database, promising users extensive information beyond mere plant identification. By acknowledging and proactively addressing the limitations identified in the literature survey, the work is set to contribute a transformative and efficient platform for Ayurvedic plant classification. In essence, the work endeavors to establish a new standard, harmonizing traditional wisdom with modern technology. Through this ongoing work, we anticipate offering a significant advancement in the exploration and dissemination of herbal medicine knowledge, ensuring the preservation of Ayurvedic botanical treasures for generations to come.

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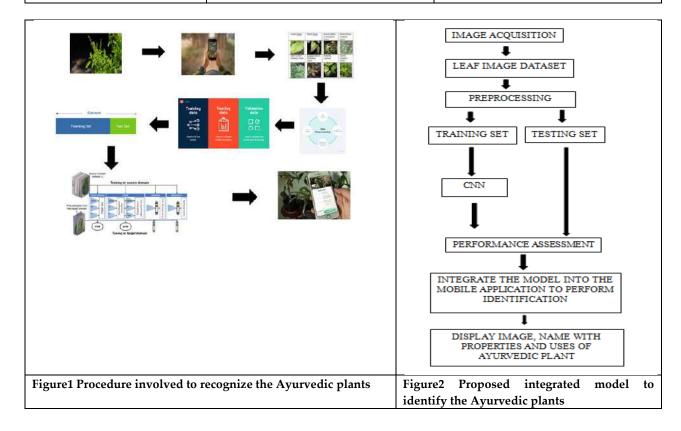
Research Paper Name	Drawback of Existing System	Proposed System
System To Detect Ayurvedic Plants and it is Medicinal Values	The proper result was not generated when the leaf images were rotated.	Designed to provide most accurate result irrespective of dimension
Identification Of Plants Using Deep Learning	Size of the dataset is limited. Samples of different classes of the plants leaf to be added	Most of the dataset has been tried to collect which are available in surrounding irrespective of classes
Worldwide Research Trends on Medicinal Plants	Discussed only about the development of medicinal plants over English medicine	Includes many other features compare to this model like identification of plant, image, information with use and benefits
Literature searches on Ayurveda	It only concentrates on Ayurveda literature search and strategy to retrieve maximum publications	It is devoid of any of the practical use that our model is proposing it is specifically designed only to search
The Significance of Ayurvedic Medicinal Plants	It discusses only on the importance, availability and caution before using the plants	Considering this information briefly our app consists a scanner to identify the plant using CNN technique and retrieve the data.
Medicinal plant classification using a convolution neural network	Identify only one plant leaf at a time and also automata plant recognition is most difficult in complex backgrounds	It is designed to identify an ayurvedic plant irrespective of complex backgrounds and also provide complete information about the plant
A mobile app for recognition	This app is only designed for plants	It is done using CNN which has highest





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of medicinal plants from Republic of Mauritius using deep learning in real time from Republic of Mauritius and it is done in MATLAB platform where they perform poorly under normal environment of 94% accuracy and the best model for plant detection and also our app contain many features with lot of benefits







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RESEARCH ARTICLE

Analysis of Traditional with Conventional RSA

S. Nirmala Devi^{1*} and M. Ganaga Durga²

¹Research Scholar, Assistant Professor, Department of Computer Applications (UG), Fatima College, Madurai, Tamil Nadu, India.

²Research Supervisor, Department of Computer Applications, Sri Meenakshi Government Arts College for Women(A), Madurai, Tamil Nadu, India.

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*Address for Correspondence

S. Nirmala Devi

Research Scholar,

Assistant Professor,

Department of Computer Applications (UG),

Fatima College,

Madurai, Tamil Nadu, India.

Email: s.nirmaladeviap@gmail.com



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ABSTRACT

During the implementation of any cryptography algorithm, it is important to consider about the computational aspect of the algorithm for the encryption and decryption process. This paper analyses the traditional and conventional RSA algorithms based on its encryption speed and decryption speed. The traditional RSA algorithm was compared with many of the other public key algorithms in various papers. But in this paper it is compared with an improved version of RSA. This improved version is proved with the help of following metrics like plain text size and key size with encryption time and decryption time. This proposed work is doing the encryption and decryption process as fast as the traditional RSA algorithm. This proposed work can be implemented for the encryption and decryption of data in strict source routing of Ad-hoc network.

Keywords: Asymmetric Encryption, Public Key Algorithm, Traditional RSA, Conventional RSA, Computational Aspects, Strict Source Routing

INTRODUCTION

Traditional RSA

This Asymmetric encryption (Public key encryption) is a form of cryptosystem in which encryption and decryption are performed using the different keys, a public key and a private key. It is also known as public-key encryption. Asymmetric encryption transforms plaintext into ciphertext using one oftwo keys and an encryption algorithm. Using the paired key and a decryption algorithm, the plaintext is recovered from the ciphertext. Asymmetric





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encryption can be used for confidentiality, authentication, or both. The most widely used public-key cryptosystem is RSA. The difficulty of attacking RSA is based on the difficulty of finding the factors of a prime number [1]. A public-key encryption scheme has six ingredients. [1]

Plaintext This is the readable message or data that is fed into the algorithm as input.

Encryption algorithm The encryption algorithm performs various transformations on the plaintext.

Public and private keys This is a pair of keys that have been selected so that if one is used for encryption, the other is used for decryption. The exact transformations performed by the algorithm depend on the public or private key that is provided as input.

Cipher text This is the scrambled message produced as output. It depends on the plaintext and the key. For a given message, two different keys will produce two different cipher texts.

Decryption algorithm This algorithm accepts the cipher text and the matching key and produces the original plaintext. The essential steps are the following.

- 1. Each user generates a pair of keys to be used for the encryption and decryption of messages[1].
- 2. Each user places one of the two keys in a public register or other accessible file. This is the public key. The companion key is kept private. Each user maintains a collection of public keys obtained from others.
- 3. If sender A wishes to send a confidential message to receiver B, sender A encrypts the message using B's public key (Fig. 1).
- 4. When B receives the message, he/she decrypts it using his/her own private key. No other recipient can decrypt the message because only B knows his/her private key.
- 5. If encryption is done using sender's private, then the receiver will decrypt it using A's public key (Fig. 2).

Conventional RSA

Like traditional RSA, conventional RSA also will have the components like Plain text, Encryption algorithm, Public and private keys and Cipher text. But here the difference is only in the encryption process and decryption process. After the generation of public and private keys the following steps were performed.

- 1. During the encryption process, a large integer (key) is raised to an integer.
- 2. Then the large key value is divided by 2 for reducing the computation time of the powers.
- 3. The divided value is now powered with the base which is doubled itself.
- 4. Till the power value is minimized, the process is continued and it produces the encrypted cipher text very fast.

simulation

Implementation of Traditional RSA

- 1. Choose two prime numbers, p = 5 and q = 7.
- 2. Find $n = pq = 5 \times 7 = 35$.
- 3. Again find \emptyset (n) = (p 1)(q 1) = 4 × 6 = 24.
- 4. Choose e such that e is relatively prime to $\mathcal{O}(n) = 24$ and less than $\mathcal{O}(n)$, choose e = 13. That is $gcd(e, \mathcal{O}(n)) = 1$
- 5. Define d such that d.e= 1 (mod \emptyset (n)). The correct value is d = 13, because 13 ×13 = 169 = (1 × 168) + 1.
- 6. The resulting keys are public key $PU_B = \{e,n\}$ and private key $PR_B = \{p,q,d\}[2]$.

Therefore $PU_B = \{13,35\} \& PR_B = \{5,7,13\}$

Encryption: $C=E_{PUB}(M)=M^e \mod n$ Decryption: $M=D_{PRB}(C)=C^d \mod n$

Consider a plaintext input of M= 33

Encryption: C=33¹³mod 35 C=33

This cipher text will be sent to receiver (B).

At receiver side, M=3313 mod 35

M=33





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Calculation of powers in Traditional RSA

Both encryption and decryption in RSA involve raising an integer to an integer power, mod n. It can make use of a property of modular arithmetic[6]:

 $[(a \bmod n) * (b \bmod n)] \bmod n = (a * b) \bmod n$

So it is possible to reduce intermediate results modulo n. This makes the calculation practical. Another consideration is the efficiency of exponentiation, because RSA dealing with potentially large exponents[3]. To see how efficiency might be increased, consider that it is possible to compute x^{16} . A straightforward approach requires 15 multiplications. This takes more time to compute the result[7].

Calculation of powers in Conventional RSA

To find 2 raised to the power 10, it is possible to multiply 9 times 2. But for finding 2 raised to the power very large number such as 1000000000, it is not easy to multiply 999999999 times 2 because it takes more time. Exponentiation by squaring helps to find the powers of large positive integers. The methodology behind the exponentiation by squaring is to the divide the power in half at each step.

$$f(x) = \begin{cases} x(x^2)^{\frac{n-1}{2}}, & \text{if } n \text{ is odd} \\ (x^2)^{\frac{n}{2}}, & \text{if } n \text{ is even} \end{cases} ----> \boxed{1}$$

Consider an example:

First divide the power by 2

 $3 \land 10 = (3 * 3) * (3 * 3) * (3 * 3) * (3 * 3) * (3 * 3)$

 $3^{10} = ((3^{30})^{10})$ [As per the above equation 1]

 $3 ^10 = 9 ^5$

The power is divided by 2 and base is multiplied to itself. So it is possible to write $3 ^ 10 = 9 ^ 5$.

Now, the problem is to find $9 ^5$

9 ^ 5 = 9 * 9 * 9 * 9 * 9

Here divide the power by 2. Since the power is an odd number here, it is not possible to do so. There's another way to represent 9 ^ 5 is as follows:

 $9^5 = (9^4) * 9$ [As per the above equation 1]

Now it is easy to find 9 ^ 4 and later multiple the extra 9 to the result

 $9 ^5 = (81 ^2) * 9$

When power is not divisible by 2, make power even by taking out the extra 9. Then it is already know the solution when power is divisible by 2. Divide the power by 2 and multiply the base to itself.

Now the problem is to find $(81 ^2) * 9$

 $(81 ^2) *9 = (81 *81) *9$

Finally divide the power by 2

 $(81^2) * 9 = (6561^1) * 9$ [As per the above equation 1]

Finally, the solution for $3 ^10 = (6561 ^1) 9 = 6561 9 = 59049$

Simulation results have been executed using python 3.12, Windows 64 bit OS, with Intel core i3 processor and 4 GB RAM. Also the results are analyzed based on plain text size and key size.

system study

Quality of Service

For all kind of cryptography algorithms, its performance is measured based on different metrics[4]&[5]. Performance measures of a proposed system are analyzed based on plaintext size and key size with encryption time and decryption time.





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Plaintext Size Based

To measure the quality of a conventional algorithm, traditional RSA as well as the conventional RSA is evaluated based on plain text size. For the different plain text size the time taken for encryption and the time taken for decryption is calculated.

Key Size Based

To measure the quality of a conventional algorithm, traditional RSA as well as the conventional RSA is evaluated based on key size. For the different key size the time taken for encryption and the time taken for decryption is calculated.

Table

The below table I shows the performance measure of the traditional and conventional RSA based on plain text size and encryption time in milliseconds for five different text files with size 3KB, 25KB, 75KB, 500KB and 10MB. The below table II shows the performance measure of the traditional and conventional RSA based on plain text size and decryption time in milliseconds for five different text files with size 3KB, 25KB, 75KB, 500KB and 10MB. The below table III shows the performance measure of the traditional and conventional RSA based on key size and encryption time in milliseconds for five different keys of P and Q. The below table IV shows the performance measure of the traditional and conventional RSA based on key size and decryption time in milliseconds for five different keys of P and Q.

Chart

The below Fig. 3 shows the shows the performance measure of the traditional and conventional RSA based on plain text size and encryption time in milliseconds for five different text files with size 3KB, 25KB, 75KB, 500KB and 10MB. The below Fig. 4 shows the performance measure of the traditional and conventional RSA based on plain text size and decryption time in milliseconds for five different text files with size 3KB, 25KB, 75KB, 500KB and 10MB. The below Fig. 5 shows the performance measure of the traditional and conventional RSA based on key size and encryption time in milliseconds for five different keys of P and Q. The below Fig. 6 shows the performance measure of the traditional and conventional RSA based on key size and decryption time in milliseconds for five different keys of P and Q.

RESULTS AND DISCUSSION

In this paper the traditional as well as conventional RSA algorithms performances are evaluated based on plain text size, key size and encryption and decryption time. During the analysis it is proved that conventional RSA is producing good result that is it is taking less encryption time and less decryption time compared with traditional RSA. This comparisons are done with the help of following metrics that is plain text size, key size and encryption time, decryption time.

CONCLUSION

This proposed system can be implemented in strict source routing method of Ad-hoc network for routing the packets from source to destination without changing the intermediate routers list. In this work it is also possible to encrypt and decrypt the data sent through this Ad-hoc network securely in a fast manner.

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Table 1: Encryption Time of Traditional and Conventional RSA with Respect to Plain Text Size

Plaintext Size	Encryption Time of Traditional RSA(ms)	Encryption Time of Conventional RSA(ms)
3 KB	0.0624	0.0609
25 KB	0.0624	0.0624
75KB	0.0565	0.0176
500KB	0.0625	0.0468
10MB	0.0685	0.0156

Table 2: Decryption Time of Traditional and Conventional RSA With Respect to Plain Text Size

Plaintext Size	Decryption Time of Traditional RSA(ms)	Decryption Time of Conventional RSA(ms)
3 KB	0.0469	0.0425
25 KB	0.0624	0.0468
75KB	0.0937	0.0937
500KB	0.0176	0.0156
10MB	0.0625	0.0312

Table 3: Encryption Time of Traditional and Conventional RSA with Respect to Key Size

Key Size (bits)	Encryption Time of Traditional RSA(ms)	Encryption Time of Conventional RSA(ms)
5 bits	0.0332	0.0312
10 bits	0.0781	0.0468
11 bits	0.0695	0.2500
14 bits	0.0468	0.0312
15 bits	0.0624	0.0468

Table 4: Decryption Time of Traditional and Conventional RSA with Respect to Key Size

Key Size (bits)	Decryption Time of Traditional RSA(ms)	Decryption Time of Conventional RSA(ms)
5 bits	0.0469	0.0468
10 bits	0.0624	0.0468
11 bits	0.0322	0.0156
14 bits	0.0595	0.2400



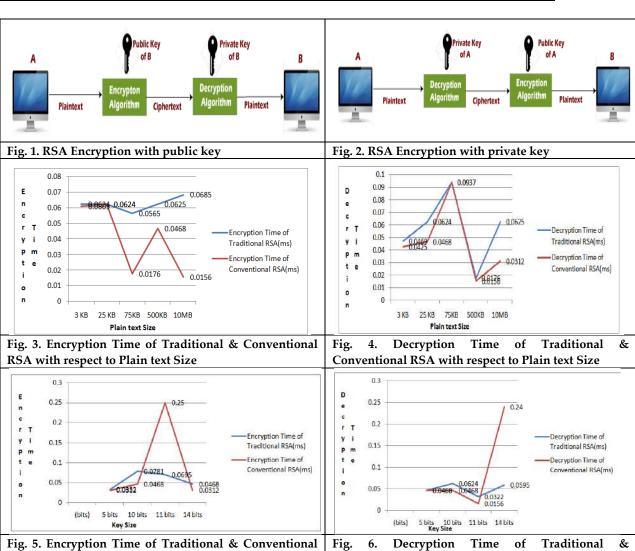
RSA with respect to Key Size



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Nirmala Devi and Ganaga Durga

15 bits 0.0468 0.0312



Conventional RSA with respect to Key Size





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RESEARCH ARTICLE

A Speech Recognition-Enabled Game using Python Integrating Twitter **Text Mining Techniques**

Simritha R1 and Tulasimala BN2

¹Student, Department of Computer Science, Mount Carmel College (Autonomous), (Affiliated to Bangalore City University), Bengaluru, Karnataka, India.

²Associate Professor, Department of Computer Science, Mount Carmel College (Autonomous), (Affiliated to Bangalore City University), Bengaluru, Karnataka, India.

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*Address for Correspondence Simritha R

Student,

Department of Computer Science, Mount Carmel College (Autonomous), (Affiliated to Bangalore City University), Bengaluru, Karnataka, India.

Email: simritha14@gmail.com



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ABSTRACT

Twitter text mining is a process of extracting valuable information from the vast amount of textual data available on the Twitter. Speech recognition process enables computers to convert spoken language into written text. In this paper, we discuss a speech recognition game that can be implemented using Python that utilizes Twitter text mining techniques and API.

Keywords: (Twitter, text mining, speech recognition, game, tweets, API)

INTRODUCTION

Speech recognition in Python is a fascinating field that involves converting spoken language into text format that computers can understand and process. It has numerous applications, including virtual assistants like Siri and Alexa, dictation software, automated customer service systems, and more. Python provides several libraries and APIs for implementing speech recognition functionality, making it accessible to developers of varying skill levels. One popular library for speech recognition in Python is Speech Recognition. It is a simple and easy-to-use library that supports multiple speech recognition engines, including Google Speech Recognition, Sphinx, and Wit.ai. With Speech Recognition, developers can quickly integrate speech recognition capabilities into their Python projects. Text mining, also known as text analytics, is the process of deriving high-quality information from text data. With the increasing availability of digital text in various forms such as documents, emails, social media posts, and more, text





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mining has become a vital tool for extracting valuable insights and knowledge from unstructured text. One interesting application of text mining is speech recognition games. These games leverage the power of natural language processing (NLP) and machine learning techniques to analyze and understand the content of tweets. By transforming the process of speech recognition into an interactive game, users can engage in a fun and educational experience while contributing to the advancement of NLP algorithms The game will provide an interactive and engaging way for users to improve their speech recognition skills while also gaining insights from Twitter data. we start by collecting tweets from Twitter using the Twitter API. The Twitter API allows us to access real-time or historical tweets based on specific search criteria or user profiles. We can filter tweets based on keywords, hashtags, user mentions, or any other relevant information. After collecting the tweets, we can preprocess them by removing noise such as URLs, special characters, stop words (commonly used words like "the," "is," etc.), and perform tasks like tokenization (splitting text into individual words), stemming (reducing words to their base form), and lemmatization (reducing words to their dictionary form). These preprocessing steps help in improving the accuracy of our speech recognition game. Once the tweets are cleaned, we will find the most frequent topics discussed in those tweets and the player will be given the five most popular texts mined using the Twitter data and will be asked to guess the word chosen randomly by the code. We will use a speech recognition library in Python, such as Speech Recognition, the user's speech input is then converted into text using Speech Recognition. The converted text is matched with the chosen random word. The player will be given three chances to guess the chosen word. By developing and deploying such a speech recognition game, we not only provide an engaging experience for users but also collect valuable data for further research and improvement of NLP algorithms. Additionally, these games can be used for educational purposes, helping users improve their speech recognition skills while having fun.

Related Work

By comparing and summarizing various methods used in speech recognition systems, the paper [1] contributes to a comprehensive understanding of the field and informs future research efforts. The authors of the research [2] state that end-to-end models represent a promising direction for automatic speech recognition, offering improved accuracy and simplified architecture. However, practical considerations such as robustness, scalability, and latency remain important factors in commercial deployment. Ongoing research efforts are focused on addressing these challenges and advancing the capabilities of E2E models for real-world applications in industry. [3] This paper provides valuable insights into the state of text mining research, highlighting its significance in big data analytics. By examining developments in methods, applications, and challenges, this review offers guidance for practitioners and researchers seeking to leverage text mining for extracting knowledge from unstructured textual data. The paper [4] provides insights into the current state and future directions of text mining applications in innovation management. By addressing methodological, conceptual, and contextual priorities, researchers can enhance the methodological rigor and richness of text mining studies in the field of innovation research.

Proposed Methodology

Text mining, also known as text data mining, is the process of extracting useful patterns, relationships, or insights from large amounts of text data. Speech recognition, on the other hand, is the ability of a computer system to recognize and transcribe human speech. In this methodology, we will be using Python to develop a text mining - speech recognition game that can extract insights from Twitter data and recognize spoken words.

Data Collection

Load Data

This step involves reading the dataset from a CSV file into a pandas Data Frame. Pandas provides efficient data structures and methods for handling tabular data, making it easy to manipulate and analyze.

Special Character Removal

Removes special characters, quotations, and leading/trailing whitespace from the text. Special characters can introduce noise and hinder analysis, so removing them helps clean the text.





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UTF-8 Encoding Handling

Handles UTF-8 encoding issues by replacing encoded characters, such as newline characters, with appropriate representations. This ensures uniformity and consistency in the text data

URL, Hashtag, and Username Removal

Removes URLs, hashtags, and usernames from the text. These elements are often irrelevant for analysis and can skew results, so removing them helps focus on meaningful content.

Lowercasing and Spelling Correction

Converts the text to lowercase to standardize it and corrects common spelling errors. Lowercasing ensures consistency in word representation, while spelling correction improves text quality.

Stopword Removal

Removes common stopwords (e.g., "the", "and") and specific words from the text. Stopwords are frequently occurring words that carry little semantic value and can be excluded to focus on more meaningful content.

Word Frequency Analysis

- 1. **Count Word Frequency:** Counts the frequency of each word in the preprocessed text to identify the most frequently occurring words. This step provides insights into the distribution of words in the dataset and helps identify key terms or topics.
- 2. **Visualize Top Words:** Visualizes the top words using a bar chart and a word cloud. The bar chart displays the frequency of the top words in a structured format, while the word cloud visually represents word frequency using font size and color, making it easier to identify prominent terms.

Speech Recognition Game.

Game Setup

Sets up the parameters for the speech recognition game, including the list of words for guessing, the number of guesses allowed, and the prompt limits. This step initializes the game environment and defines the rules.

Initialize Components

Initializes the speech recognition components, including the recognizer and microphone objects. Speech recognition libraries provide APIs for capturing and transcribing speech input, which are essential for implementing the game

Word Selection and User Prompting

Chooses a random word from the list and prompts the user to speak, initiating the game. Random word selection adds variability and excitement to the game, while user prompting encourages participation.

Speech Input Recognition

Records the user's speech input using the microphone and recognizes the transcription using the speech recognition library. This step captures the user's response and converts it into text for comparison.

Comparison and Feedback

Compares the user's guessed word with the chosen word and provides feedback to the user based on the correctness of the guess. Feedback can include messages indicating whether the guess was correct or incorrect.

Multiple Attempts and Game Termination

Allows the user multiple attempts to guess the word within the specified number of guesses. The game terminates based on whether the user correctly guesses the word or exhausts all attempts. This step concludes the game and provides closure to the user's interaction.





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RESULTS AND DISCUSSIONS

The results underscored the importance of data preprocessing in text mining projects and highlighted the potential applications of text analysis techniques in various domains. The initial phase of the project involved preprocessing the Twitter data extracted from the 'full-corpus.csv' file. The preprocessing steps included:

- 1. Removal of leading and trailing whitespaces.
- 2. Replacement of newline characters with spaces.
- 3. Substitution of UTF-8 encoding with corresponding characters such as single quotes and ellipses.
- 4. Elimination of all other UTF-8 encoding patterns.
- 5. Truncation of text after any 'https:' occurrences.
- 6. Removal of text before colon (':') occurrences.
- 7. Omission of hashtags and usernames.
- 8. Elimination of special symbols such as punctuation marks and currency symbols.
- 9. These preprocessing steps aimed to standardize the textual data and prepare it for further analysis.

Word Frequency Analysis

Following data preprocessing, we conducted word frequency analysis to identify the most common words in the dataset. The analysis revealed several frequently occurring words, including 'legend,' 'harambe,' and 'rip.' The distribution of word frequencies was visualized using a bar chart, which displayed the top 50 most common words and their respective counts.

Word Cloud Visualization

To provide a visual representation of the text data, we generated a word cloud using the 'WordCloud' library in Python. The word cloud depicted the most prevalent words in the dataset, with larger font sizes indicating higher frequencies. The visualization offered insights into the dominant themes and topics present in the Twitter data.

Speech Recognition Game

As a practical application of the processed text data, we implemented a speech recognition game using the 'speech_recognition' library. The game involved randomly selecting a word from the dataset and prompting the user to guess the word by speaking into the microphone. The program then recognized the user's speech and determined whether the guess was correct.

CONCLUSION

The results of the data preprocessing phase demonstrated the effectiveness of the implemented cleaning techniques in standardizing the textual data. The word frequency analysis provided valuable insights into the prevalent topics and discussions on Twitter during the data collection period. Additionally, the word cloud visualization offered a visually appealing representation of the most common words, facilitating easier interpretation of the dataset. The speech recognition game showcased the practical utility of natural language processing techniques in interactive applications. By integrating speech recognition functionality with the processed text data, we demonstrated a real-world application of text mining methodologies. Overall, the results underscored the importance of data preprocessing in text mining projects and highlighted the potential applications of text analysis techniques in various domains.

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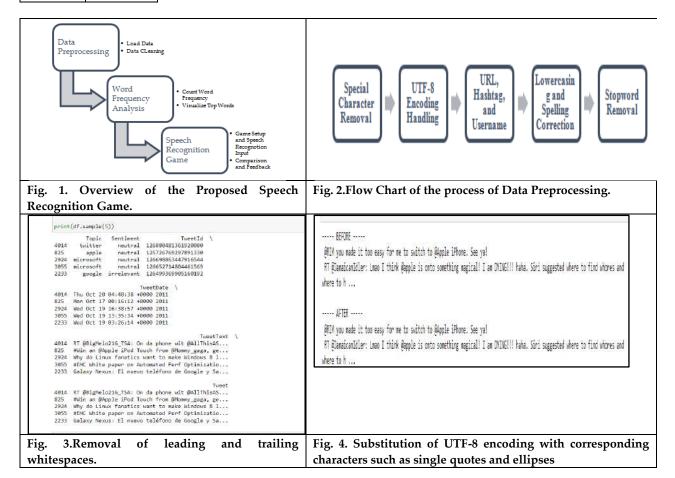


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Table 1. Count of Word and Frequency

Word	Frequency
new	239
cream	173
sandwich	170
android	168
ice	163
google	159
nexus	156







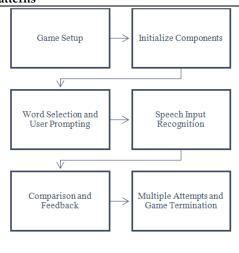
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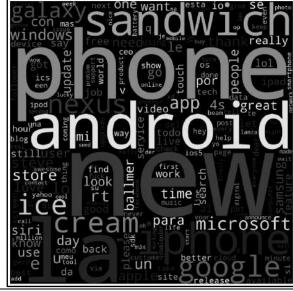


Fig. 5. Elimination of all other UTF-8 encoding patterns

remove words = ['3', 'long', 'years', 'year', 'goes', 'cant', 'today', '\'', 'ago', 'later', three', fucking', 'nuisee', 'happon', much', 'fishrich', 'shit', 'tweet', 'in', 'follon', 'lifete', 'think', well', 'will', 'going', 'tuither', 'min't', 'right', 'mon', '\\ 'all', 'seess', 'inreme', 'min', 'joe', 'gaee, 'naee', 'ell', 'wall' 'gst', 'ye', 'yet', 'charity', 'telethon', 'dont', 'said', 'tevnofs, 'make', 'see, 'internet', 'amp', 'en', 'est', 'el', 'hn', 'dm', 'na', 'de', 'que']

Fig. 6. Removal of common stopwords





7. Working process Fig. the Speech of Recognition-enabled game

Fig. 8.Word Cloud of the most frequently occurring words in the tweets

I'm thinking of one of these words: new, cream, sandwich, android, ice You have 3 tries to guess which one.

Guess 1. Speak! You said: Android Incorrect. Try again.

Guess 2. Speak! You said: sandwich Correct! You win!

I'm thinking of one of these words: new, cream, sandwich, android, ice You have 3 tries to guess which one. Guess 1. Speak! You said: hello Incorrect. Try again. Guess 2. Speak! I didn't catch that. What did you say? Guess 2. Speak! I didn't catch that. What did you say? Guess 2. Speak! You said: cream Incorrect. Try again. Guess 3. Speak! You said: Android Sorry, you lose! I was thinking of 'sandwich'.

Fig. 9.Speech Recognition game where the user guessed the word - sandwich correctly

Fig. 10.Full working of the game where the user fails to guess the word after 3 tries





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RESEARCH ARTICLE

IoT - Powered Bio sensors Lead the way in Environmental Sensing in the **Digital Age**

Sumanth S1* and Siddarama S2

¹Associate Professor, Department of Computer Science and Applications, Government College for Women, Kolar – 563 101, Karnataka, India.

²Head, Department of Computer Science and Applications, Sri Bhagawan Mahaveer Jain College, Robertsonpet, K.G.F. – 563 122, Karnataka, India.

Received: 20 Dec 2023 Revised: 29 Jan 2024 Accepted: 03 Mar 2024

*Address for Correspondence

Sumanth S

Associate Professor,

Department of Computer Science and Applications,

Government College for Women,

Kolar – 563 101, Karnataka, India.

Email: sumanth81s@gmail.com/dr.sumanth.s.81@gmail.com



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ABSTRACT

In today's age of digital transformation, the convergence of environmental sensing and IoT has enabled previously inconceivable advancements in ecological monitoring and management. This research looks into how Internet of Things (IoT) technology and biosensors can be used to create an all-encompassing system for real-time environmental monitoring. IoT-powered biosensors are strategically positioned throughout ecosystems to monitor and report on air and water quality, pollution levels, biodiversity, and other factors. This data is used to create early warning systems for environmental hazards, to encourage biodiversity conservation, to ensure industrial compliance with legislation, and to steer legislators toward more evidence-based policies. The system will delve into the workings of these biosensors in this study, emphasizing their relevance in safeguarding ecosystems, keeping the public healthy, and paving the way for future research. Furthermore, it addresses sensor calibration, data management, privacy, and efficiency issues. This study employs an interdisciplinary approach combining biology, engineering, and data science to explain how IoT-powered biosensors are ushering in a more sustainable and environmentally conscious future by vastly improving our ability to monitor, understand, and proactively address environmental challenges.

Keywords: Environmental Monitoring, Biodiversity, Pollution Levels, Water Quality, Internet of Things (IoT).





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INTRODUCTION

In an age marked by tremendous technical advancements, humanity is at a crossroads in the quest to address pressing environmental challenges. Human-caused climate change, air and water pollution, biodiversity loss, and habitat degradation pose major threats to ecosystem health and human and nonhuman animal existence. Confronting these challenges necessitates the development of creative ways for improving environmental monitoring and management. The IoT and environmental biosensors are converging to become a game-changing factor in this effort [1]. The IoT revolution, defined by the internet's penetration into previously unconnected spaces, has ushered in a new era of data collection, analysis, and decision-making in many fields. Because of its networked nature, it has the potential to revolutionize environmental research, where manual data gathering and sporadic sampling have long been the norm for keeping tabs on environmental factors in real time. When IoT devices are coupled with biosensors designed to interact with and detect biological factors, a whole new level of continuous, high-resolution data collection from the natural world becomes possible. These IoT-enabled biosensors, also known as environmental or biologically inspired sensors, keep an eye on the environment and report any real-time changes [2]. This study investigates the promising new direction in environmental monitoring and management by combining IoT technologies with biosensors. Our research digs into this merger's fundamental ideas, approaches, applications, and ramifications, illuminating how it can radically alter how we understand and engage with the natural world [3]. The IoT, combined with biosensors, has revolutionary potential in many environmental research and sustainability areas. Simply put, this convergence improves our ability to gather timely and precise information on environmental conditions, allowing for more effective, preemptive measures to be taken in the face of new dangers and the development of policies based on solid empirical evidence [4].

The infrequent nature of the conventional environmental monitoring techniques has meant that this has been inadequate. Periodic data gathering is used by many research institutions, government organizations, and companies, yet this method leaves important knowledge gaps about environmental dynamics. Biosensors powered by the IoT can monitor environmental variables in real time. This capacity becomes very important when dealing with environmental emergencies like natural disasters or industrial mishaps [5]. Early warning systems that can identify environmental abnormalities and dangerous circumstances in real time can potentially reduce the severity of catastrophes and safeguard human health. For instance, air quality monitors may send out warnings when unhealthy air is in the area, giving people time to take precautions and giving hospitals time to prepare for an uptick in respiratory infections [6]. Loss of habitat, warming temperatures, and pollution pose serious biodiversity problems. Biosensors powered by the IoT may help biodiversity conservation by tracking animal activity in the wild. Conservation efforts and methods may be improved with the knowledge gained by researchers on migratory patterns, reproductive habits, and responses to environmental changes [7]. Several sectors contribute to pollution and trash that might be harmful to ecosystems.

To ensure that businesses comply with environmental rules, biosensors powered by the IoT may be used to track pollutants in real time. This not only aids in environmental defense but also encourages environmentally responsible manufacturing [8]. Advanced data management and analytics solutions are required due to the enormous data output from IoT-driven biosensors. However, when properly used, this information equips decision-makers with knowledge of environmental patterns over the long term and guides the creation of policies meant to solve environmental concerns [9]. Researchers in environmental science may get a wealth of useful information from biosensors powered by the IoT. This plethora of data makes it easier to comprehend ecological processes, aids in detecting new research problems and quickens the rate of scientific progress. Connected biosensors powered by the IoT have the potential to raise awareness and inform the public about environmental concerns in real-time. This technology may be used by citizen science projects to get more people involved in environmental monitoring and study. This study, in summary, sets out on a voyage through the world of IoT-driven biosensors in environmental monitoring. It is a journey packed with promise, obstacles, and the possibility of profound change [10].





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Related Work

In this day and age, it is critical to perform frequent environmental monitoring to assess the state of ecosystems, protect public health, and influence policy decisions. Recently, IoT-based biosensors and platforms such as Raspberry Pi have been developed, enabling for more precise, data-driven environmental monitoring. [11]. Biosensors based on the IoT have become an effective method of monitoring several environmental factors. Changes in air and water quality, pollution levels, and biodiversity indicators may all be detected with the help of these sensors, which use a wide range of identification processes (including chemical, biological, and auditory). These sensors' capacity for continuous, distant data collecting provides a detailed picture of the surrounding environment, allowing for the rapid identification of abnormalities and potential dangers [12]. Adding Raspberry Pi to IoT biosensors improves their performance. The Raspberry Pi acts as the network's brain, handling and transferring information in near-real time. To get more people involved in environmental monitoring, it also helps to provide user-friendly interfaces for data visualization and accessibility [13].

IoT-based biosensors have found widespread use in air quality monitoring. Particulate matter (PM2.5, PM10), carbon dioxide (CO2), nitrous oxides (NOx), volatile organic compounds (VOCs), and volatile gases may all be detected by these sensors. To determine the effects of air pollution on respiratory health and adopt preventative actions, constant air quality monitoring is essential for public health [14]. The safety of drinking water and the preservation of aquatic ecosystems highlight the need to monitor water quality. Heavy metals, organic pollutants, and changes in water pH and turbidity are only some of the contaminants that may be detected by biosensors built on the IoT. This information is helpful to better manage water resources and reduce pollution [15]. Using live microorganisms as sensors to detect pollution is a novel strategy. These biosensors change their behavior or metabolic activity when exposed to various contaminants. Timely actions to reduce pollution's negative effects and save aquatic life are made possible through quick responses. The existence, behavior, and population dynamics of species in different habitats may be better understood via biodiversity monitoring using auditory and visual recognition technology. This information is useful for biodiversity conservation and understanding climate change's ecological effects. Using Raspberry Pi and other IoT-based biosensors is an exciting step forward in environmental monitoring. To better safeguard public health and further scientific understanding, stakeholders may benefit from the convergence of data, biology, and technology. Creating such monitoring systems may fuel more effective environmental conservation and sustainability campaigns.

PROPOSED METHODOLOGY

Work model

Our long-term goal in developing IoT-based biosensors for environmental monitoring is to create a network of sensors that can continuously monitor their surroundings and respond to any changes they notice. Our strategy takes into account the creation and deployment of these biosensors, as well as their impact on public health and environmental safety across their entire lifecycle. Carbon monoxide (CO), nitrogen oxides (NOx), sulfur dioxide (SO2), volatile organic compounds (VOCs), and particulate matter (PM2.5, PM10) are just some of the gases and particles we hope to one day be able to detect and measure with our sensors. Gas-sensitive materials and electrochemical transducers are used in these detectors. The molecular recognition concept is at the heart of the air quality sensor's operation, with its constituent parts interacting selectively with the gases of interest. When these reactions occur, an electrical current or potential difference is created on the sensor's surface. The concentration of a target gas may be used to accurately calibrate these variations, allowing for very accurate environmental monitoring. The system developed a pollution sensor that mimics the function of naturally occurring microbes as a holistic approach to cleaning up polluted water sources. This sensor uses live microorganisms that provide distinctive signals in response to individual contaminants. There is clear evidence that these microbes' metabolic processes or behaviors alter when exposed to polluted water. Our pollution sensor detects the presence of different contaminants and provides crucial insights into water pollution levels by monitoring these biological reactions. The integration of sound sensors and picture identification software forms the backbone of our biodiversity sensor. This sensor





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exemplifies how far biosensing technology has come in recent years. While image recognition algorithms evaluate visual data, acoustic sensors record ambient noise. The biodiversity sensor is a device that, when placed in natural environments, takes pictures of plants and animals and records their unique audio signatures. It may learn more about the existence of species, their activities, and the dynamics of their populations by evaluating these signals and photos, which in turn helps with conservation efforts. Our biosensors depend on wireless communication to deliver data on air and water quality, pollution levels, and markers of biodiversity once we have acquired this information. Our sensors transmit information to a centralized database or cloud-based platform through low-power, wide-area networks (LPWANs) or cellular networks. This wireless connection guarantees a steady data supply, allowing for continuous, real-time tracking. When it comes to the IoT, data security is of the utmost importance. It uses strong encryption and authentication methods to keep information secure and private while in transit. This guarantees the privacy and integrity of the information our biosensors capture as it travels from device to database. Our internetenabled biosensors upload their data to a central database on the cloud. This database serves as a nerve center, collecting and organizing sensor information across various environments. This architecture ensures the security and availability of data. While raw sensor data is rich, it frequently needs significant processing before being used to conclude. This is when our data analytics algorithms come into play. These algorithms analyze data in real-time, spotting trends, outliers, and other irregularities that may indicate ecological danger. Our biosensors give decisionmaking help via continuous data analysis. It is crucial to make data accessible and easy to grasp for a large audience. It provides intuitive dashboards and other visual aids for our clients to help them do this.

Visualizing sensor data in charts, graphs, and maps, these tools help academics, policymakers, and the general public understand how the environment changes over time. Each monitored metric has predetermined thresholds that form the basis of our early warning systems. The scientific community and environmental regulators work together to set these limits. These are used as benchmarks against which sensor data in real-time may be evaluated. Our early warning systems are activated when sensor readings cross specified limits. Potential environmental concerns are identified, and the appropriate parties are notified through alerts. These warnings are very helpful for making prompt decisions and countering ecological dangers. Our guiding idea materializes as a sophisticated network of IoT-based biosensors that track real-time environmental metrics like pollution levels and biodiversity indices. The data is continually uploaded to a hub where it may be examined in real-time and shared with many people. It can respond quickly to new environmental risks by establishing clear thresholds and implementing early warning systems. It helps to preserve ecosystems and biodiversity, reduce pollution, and advance sustainability by monitoring and reacting to environmental changes and threats in real-time. Monitoring air and water quality in real time is important for public health because it gives local populations the information they need to take precautions against potential threats. Our biosensors create data useful for scientific investigation, enhancing our comprehension of environmental processes and adding to the corpus of knowledge. Using this information, policymakers and environmental agencies may make data-driven decisions that better protect the environment. Our IoT-based biosensors are designed to protect the environment, advancing science and improving public health as their guiding principles. The future of Earth and all its inhabitants may be made more secure and sustainable if It keeps an eye on and adapts to environmental changes in real-time.

Description

The gases and particles in the air directly affect human health and the environment, and our air quality sensor is a highly developed gadget designed to detect all of these contaminants. This sensor is based on molecular identification and works with gas-sensitive materials and electrochemical transducers. CO2, nitrogen oxides (NOx), sulfur dioxide (SO2), volatile organic compounds (VOCs), and particulate matter (PM2.5, PM10) all set off a targeted chemical reaction in the sensor. This reaction results in surface-level electrical current or potential changes, allowing for accurate air quality assessments. It allows us to track and react to changes in air quality and pollution levels, making our communities safer and more hospitable. The model is shown in Figure 1. The inherent recognition systems present in living creatures served as inspiration for our biomimetic water quality sensor. Heavy metals, organic pollutants, pH changes, and turbidity levels are only some contaminants that this sensor may detect. It is based on molecular recognition theory, which states that certain "recognition elements" may interact preferentially





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with "target substances" in water. The sensor detects biological events triggered by this contact as changes in its electrical characteristics. These shifts may be measured, allowing for precise evaluations of water quality. This sensor will help us protect freshwater ecosystems, keep an eye on the purity of our drinking water, and react quickly to any new problems we may have with water quality. Our pollution sensor is a novel approach to the problem of water pollution. This sensor takes its cues from the capacities of naturally occurring microbes and uses them to detect the presence of very particular contaminants in water. These bacteria modify their metabolic processes or behavior in unique ways when exposed to polluted water. The sensor records these biological reactions, illuminating the prevalence and severity of pollution in water systems. The system can safeguard aquatic ecosystems and encourage regulatory steps to lower pollutant levels in water supplies if It knows where the pollution comes from and how often it occurs. Our biodiversity sensor fully displays a combination of sound sensors and picture recognition software. This sensor may be used in various environments to keep tabs on species' existence, behavior, and population dynamics. While image recognition algorithms evaluate visual data, acoustic sensors record ambient noise. The biodiversity sensor is placed in natural environments to record the sounds and photos of many animals. Examining these signs and pictures, the system learns about ecosystems' richness and vitality.

The data gathered by this sensor will help scientists, conservationists, and legislators choose where to focus their efforts to protect biodiversity. Raspberry Pi is a local data storage system that may be used to keep track of environmental data over time and to do long-term trend analysis. It may organize sensor data for later retrieval and use as a reference. Data redundancy and preservation may also be ensured by storing more information on the cloud or external storage devices. Since Raspberry Pi is so adaptable, it can be easily combined with weather monitoring systems like weather stations, remote webcams, and weather prediction services. Raspberry Pi improves the environmental monitoring ecosystem by compiling data from various sensors to provide a complete picture of the state of the world around us. Raspberry Pi is the brains of our IoT biosensors for monitoring the surrounding environment. It manages sensor data in real-time, allowing for advanced analytics, early warning systems, and intuitive user interfaces. The addition of Raspberry Pi improves the capabilities of our biosensors, making them a potent resource for data-driven decision-making, ecological safeguarding, and citizen participation in the fight for a more sustainable and resilient world.

RESULTS AND DISCUSSIONS

The technology has been extensively researched, with these sensors used in a range of ecosystems to improve environmental monitoring utilizing IoT-based biosensors coupled to Raspberry Pi. Our environmental monitoring effort generated the data presented in this section, which was rigorously examined to assess its importance. Our air quality monitors found significant air contaminants with surprising precision. Throughout the research, our sensors collected constant data on numerous gasses and particles. The sensors were capable of monitoring CO2 levels both inside and outside. CO2 levels vary according to human activity, ventilation, and environmental variables. The sensors precisely detected NOx and SO2 levels, revealing higher concentrations in urban and industrial areas. The device measured PM2.5 and PM10 concentrations constantly to assess the effects of air quality on respiratory health. There was a link between high PM levels and unfavorable health consequences, emphasizing the importance of addressing air quality issues. Many pollutants were detected by our biomimetic recognition-inspired water quality monitors. Some water sources were determined to have dangerous amounts of heavy metals such as lead, mercury, and cadmium, spurring more research into the core causes of pollution and potential solutions. Organic pollutants including pesticides and industrial chemicals were found in water samples by the sensors, highlighting the need for tougher controls and more sustainable farming practices. Changes in pH and turbidity have been recorded, which may have ramifications for aquatic ecosystems and necessitate specific management techniques. Using living bacteria as sensors, the system was able to gather critical information regarding contamination levels in water systems. The sensors monitored variations in microbial activity in order to identify pollution sources and direct cleanup activities. It was discovered that in reaction to pollution incidents, prompt steps were conducted, allowing for timely interventions to decrease ecological harm and protect aquatic life. Using sound and image recognition technologies,





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our biodiversity sensors could accurately record species richness in their natural habitats. The sensors recorded the existence of several species, expanding our knowledge of the biodiversity in the area. Acoustic data showed distinct behavioral patterns of several species, providing insight into these hitherto unknown organisms' ecological functions and interconnections. Our sensors revealed information on population dynamics and the effects of environmental changes on biodiversity by monitoring long-term shifts in species abundance. Our sensors' ability to perform continuous monitoring enables them to identify environmental changes and potential dangers at an earlier stage. Data with a high resolution about the quality of the air and water, the levels of pollutants, and the biodiversity of an area might prompt actions, therefore averting or minimizing ecological harm and protecting human health. The example sensor data in Tables 1 and 2 comes from various environmental monitoring sensors and has been split up for easy reading. Environmental indicators include carbon monoxide and nitrogen oxides (NOx), lead and pH in water, and bird species diversity. The data consists of several dimensions: measurement units, sensor locations, timestamps, and values. Please keep in mind that the information below is only indicative.

The vast amounts of gathered data allow academics, environmental agencies, and lawmakers to make choices based on accurate information. Analysis of data collected in real-time makes it possible to implement adaptive management solutions, which in turn leads to environmental policies and actions that are more successful. Using our sensors encourages public participation in initiatives to monitor the environment. User-friendly interfaces hosted on Raspberry Pi make it possible for citizen scientists and local communities to access data collected by sensors. Through these interactions, knowledge is raised, environmental stewardship is encouraged, and collective action to safeguard natural resources is encouraged. The information collected by our biosensors contributes to advancing scientific knowledge and comprehending the processes that occur in the environment. Researchers can use this data to investigate the impacts of pollution, climate change, and human activity on ecosystems over the longer term. Our sensors show the necessity for management strategies tailored to individual ecosystems. The identification of diverse pollution sources and shifts in the pH of the water, for instance, highlights the need to develop individualized plans for various ecosystems. Monitoring the air quality in real-time, especially the levels of particulate matter and gaseous pollutants directly affects the general population's health. Informing the public and authorities about the risks of exposure and enabling timely preventive action is a potential outcome. Our IoT biosensors integrated with Raspberry Pi are revolutionary for tracking environmental conditions. In pursuing a more resilient and sustainable global future, our sensor data has great potential for enhancing decision-making, public involvement, and scientific development. This system protects our world and its people using data, biology, and technology.

CONCLUSION

Integrating IoT-based biosensors with Raspberry Pi to improve environmental monitoring offered significant insights as well as the potential for disruptive change. The outcomes of this study demonstrate how beneficial such sensors could be for real-time monitoring of environmental parameters such as air and water quality, pollution levels, and biodiversity markers. Sensor data enables us to make informed decisions, which speeds up the detection and response to environmental changes and dangers. Furthermore, the availability of real-time sensor data made accessible via user-friendly interfaces benefits both public participation and scientific knowledge advancement. The integration of technology, biology, and data will become increasingly crucial in preserving natural resources, defending public health, and protecting the environment as time goes on. Our journey will continue to commit to improving these monitoring systems, expanding the spatial coverage, and encouraging collaboration among many stakeholders. Finally, this effort contributes to the larger goal of creating a sustainable and resilient world for current and future generations.





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Table 1: Data

Parameter	Measurement Unit	Location	Time Stamp	Value
Air Quality (CO2)	ppm (parts per million) Indoo		2023-09-01 09:00	400
Air Quality (CO2)	ppm	Indoor	2023-09-01 10:00	410
Air Quality (CO2)	ppm	Indoor	2023-09-01 11:00	415
Air Quality (NOx)	ppm	Outdoor	2023-09-01 09:00	20
Air Quality (NOx)	ppm	Outdoor	2023-09-01 10:00	22



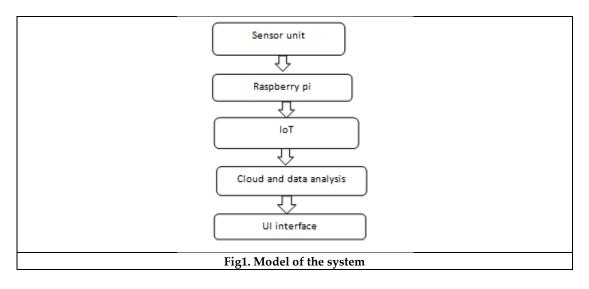


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Air Quality (NOx)	ppm	Outdoor	2023-09-01 11:00	25
Water Quality (Lead)	μg/L (micrograms per liter)	River	2023-09-01 09:00	5
Water Quality (Lead)	μg/L	River	2023-09-01 10:00	6
Water Quality (Lead)	μg/L	River	2023-09-01 11:00	7

Table 2: Diversity Data

Parameter	Measurement Unit	Location	Time Stamp	Value
Water Quality (pH)	рН	Lake	2023-09-01 09:00	7.2
Water Quality (pH)	рН	Lake	2023-09-01 10:00	7.1
Water Quality (pH)	рН	Lake	2023-09-01 11:00	7.0
Biodiversity (Birds)	Count	Forest	2023-09-01 09:00	25
Biodiversity (Birds)	Count	Forest	2023-09-01 10:00	30
Biodiversity (Birds)	Count	Forest	2023-09-01 11:00	27







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RESEARCH ARTICLE

Diagnosis of Grape Leaf Disease using CNN

Kiruthika S1*, Suriya K2 and Pooja S2

¹Assistant Professor, Department of Information Technology, Sri Ramakrishna College of Arts & Science, Coimbatore, Tamil Nadu, India.

²II MSc IT, Department of Information Technology, Sri Ramakrishna College of Arts & Science, Coimbatore, Tamil Nadu, India.

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*Address for Correspondence

Kiruthika S

Assistant Professor, Department of Information Technology, Sri Ramakrishna College of Arts & Science, Coimbatore, Tamil Nadu, India.

Email: kiruthika@srcas.ac.in

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ABSTRACT

Grape leaf diseases are a major problem for crops used in agriculture, leading to significant losses in terms of money and environmental damage. Early identification and control are essential to curb the spread of leaf diseases and lessen their detrimental impact on crop quality and production. However, accurately diagnosing leaf diseases and selecting the appropriate pesticide treatments can be challenging undertakings for farmers and agricultural professionals. The variety of diseases that can negatively impact vineyard productivity and overall stability in grapevines is a significant issue for the grape business. Timely identification and precise diagnosis are essential for minimizing monetary losses and safeguarding the vineyard environment. The research focuses on classifying images of grape leaves into several disease categories, providing a significant advancement towards sustainable vineyards and worldwide.

Keywords: Grape leaf, Deep Learning, CNN, Mobilenetv2

INTRODUCTION

A highly prized fruit in the agricultural industry, grapes can suffer from a number of diseases that negatively impact both its quality and productivity. Detecting and classifying diseases effectively is crucial to handling these problems. This work presents a novel machine learning method for disease classification in grape leaves [1]. The make use of a broad collection of high-resolution photos that include samples of both leaves in good condition and leaves afflicted with prevalent grape diseases like black rot, black measles, and downy mildew. It creates reliable and accurate illness





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classification models by applying cutting-edge machine learning algorithms, such as Transfer Learning and ensemble methods, and sophisticated pre-processing techniques. The outcomes demonstrate these models' potential for useful application in grape vineyards and other agricultural contexts. The results of this study will have a big impact on the grape growing sector. Grape growers and vineyard managers can take use of automated and dependable disease diagnosis with the created machine learning models, which will allow them to quickly contain disease outbreaks and protect crop health.

IIRELATED WORK

To avoid large crop losses, crop diseases must be identified early and managed. Deep convolutional neural networks (CNN) were suggested by Nagi (2022) as a method for classifying grapevine leaf species. The CNN model can be utilised as a tool for automated species identification because it demonstrated great accuracy in distinguishing between several grapevine leaf species. In their 2020 study, Ghoury et al. sought to accomplish tasks with a transfer learning technique that uses pre-trained deep learning models, such as Faster R-CNN Inception v2 and Single SSD_Mobile Net v1, to discriminate between sick and healthy grapes and grape leaves. There were 136 healthy photos and 124 sick images in the image dataset that was used. Ghosh (2020) begins by segmenting the first affected area, extracting texture and colour information, then resizing them to the desired pixel. Data collection, image processing, and image segmentation were the processes that were put into practice. After that, these pictures go through a few pre-processing steps, like image scaling and image smoothing. After preprocessing the data, features in leaf images are extracted using techniques like RGB (color-based features). The names are then changed to categories for model compatibility, based on the photos' matching class. Eighty percent of the training data are used to model the classifier using neural network techniques. The model is validated using the remaining 20 percent of the data. A 92% accuracy rate was achieved with this strategy. Liu Y (2020) describes a convolutional neural network (CNN) based approach for classifying grape varieties based on leaf pictures. The scientists take information from leaf pictures and feed them into a CNN model to categorise different grape varietals.

METHODOLOGY

CONVOLUTIONAL NEURAL NETWORK

One kind of neural network used for processing and categorising images is called a convolutional neural network (CNN) [3]. They are made up of multiple layers, such as a fully connected layer, a pooling layer, and a convolutional layer. The network extracts elements like edges, corners, and textures from the input image in the convolutional layer by applying a series of filters on it.

Mobile Net

The initial architecture, known as Mobile Net, was unveiled in 2017. Its main objective was to create a neural network architecture that was small enough to operate well on mobile devices and still produce accurate results.

MobileNetV2

A neural network architecture called MobileNetV2 was created for effective deep learning on mobile and embedded platforms. It expands on the original Mobile Net architecture with the goal of increasing efficiency and accuracy

DATA COLLECTION & PRE-PROCESSING INTRODUCTION

Any research endeavour, including the creation of an AI-based framework for the identification of leaf diseases, must include data collection. We used an existing dataset that we obtained from Kaggle for research. In order to acquire experimental data, an artificial environment simulating various crop disease conditions is created. In order to gather experimental data for this study, crops will be purposefully infected with a variety of illnesses, and the affected leaves will subsequently be photographed. The accuracy of the suggested based framework in diagnosing leaf diseases will be trained and validated using this data.





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SPLITTING THE DATASET

The process of building a dataset for machine learning involves first using augmentation techniques [14], and then dividing it into three sections: training, validation, and testing sets. With 70% of the dataset being used for training, 20% for validation, and 10% for testing. The augmented photos in the training set are used to train the machine learning model, while the validation set is used to track the model's performance and make necessary parameter adjustments to increase accuracy. To make sure the model can generalise successfully and perform accurately on new data, the testing set is used to assess the trained model's ultimate performance on fresh, unknown data.

IMAGE PROCESSING

Using CNN methods and techniques, image processing in real-time dataset development entails extracting features and valuable information from images. The procedure includes capturing the image, pre-processing it, extracting features, and classifying the results. It may be used to detect crop illnesses, track crop development and production, and offer timely crop management advice. Farmers may enhance overall crop performance, maximise yield and quality, and make timely decisions about crop management by evaluating crop health and growth. A more profitable and sustainable agricultural business can result from the identification and mitigation of risks related to crop production through the use of image processing.

CLASSIFICATION OF THE LEAF DIASEASE INTRODUCTION

Numerous diseases that damage grapevine leaves have been identified, which poses a serious challenge to grape production. The grape industry suffers large financial losses as a result of three common grape leaf pests and diseases: black rot, esca (black measles), and leaf blight (Isariopsis leaf spot). in order to stop the spread of illness and guarantee the grape industry's continuous expansion. Accurate detection and identification of illnesses affecting grape leaves are crucial. Figure: Grapevine leaves afflicted by illness Convolutional neural networks (CNNs) have demonstrated remarkable efficacy in image classification tasks; nevertheless, training them from the beginning necessitates substantial volumes of labelled data and can be computationally demanding. By using pre-trained CNN models that have picked up pertinent features from massive datasets, transfer learning has become a potent method for overcoming these difficulties. Transfer learning is the process of refining a pre-trained model—like a CNN—on a fresh dataset tailored to the current classification task after it has been trained on a sizable dataset, like Image Net. The pre-trained model's layer weights are changed during this fine-tuning procedure, which also entails training the model on the new dataset to enable it to acquire high classification accuracy for photos. Transfer learning greatly reduces the amount of data and computation needed to train an efficient image classification model by utilising the features learned by the pre-trained model.

TRANSFER LEANING

A pre-trained model is used as the basis for a new machine learning challenge in a process known as transfer learning [4]. A sizable dataset was used to train the pre-trained model, which taught it to identify pertinent features. Then, using a smaller dataset, these features can be used to tackle a new, related job. Transfer learning can increase the accuracy of the new model while saving time and computational resources by beginning with a pre-trained model. Back propagation, a technique, allows the pre-trained model to have its weights adjusted for the fresh dataset, allowing for further refinement. Natural language processing, picture categorization, and other machine learning fields have all seen a rise in the use of transfer learning.

MODEL 1 – MOBILENET MODEL

Google researchers created Mobile Net, a flexible convolutional neural network architecture designed for effective deep learning on embedded and mobile devices. Depth-wise separable convolutions, which Mobile Net uses instead of typical convolutional neural networks, greatly lessen the processing load. By making this design decision, Mobile Net is able to minimise the size and processing requirements of the model and still give attractive performance. Mobile Net is particularly well-suited for picture categorization, object recognition, and other computer vision applications on devices with limited resources because of its lightweight design. The fact that Mobile Net can





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perform at the cutting edge of picture classification tasks while using less processing power than other deep learning models highlights the efficiency of this model. Its usefulness goes beyond mobile and embedded devices, as it can also be a vital instrument for real-time computer vision applications. This highlights the need of effective model construction in the context of artificial intelligence and deep learning.

MobileNetv2

Researchers at Google created the novel convolutional neural network architecture known as MobileNetV2, which is similar to Mobile Net. It is a noteworthy development in the field of deep learning, particularly in terms of the effective application of neural networks on embedded and mobile devices. The goal of MobileNetV2 was to improve the trade-off between model size and The adaptability of MobileNetV2 to various model configurations through the use of hyper parameters such as the width multiplier and resolution multiplier is one of its noteworthy features. By adjusting the width multiplier, users can balance the model's number of channels between accuracy and size. One example of the ongoing efforts to create effective neural networks is MobileNetV2. Convolutional neural networks (CNNs) of the Mobile Net class were made available as open-source software by Google, making them a great place to start for developing really small and incredibly quick classifiers. The number of multiply-accumulates (MACs), a measure of the number of fused multiplication and addition operations, determines the network's speed and power consumption. A family of Tensor Flow computer vision models called Mobile Nets is focused on mobile devices and is intended to optimize accuracy while taking into account the limited resources of embeddedoron-device applications. The number of multiply-accumulates (MACs), a measure of the number of fused multiplication and addition operations, determines the network's speed and power consumption. A family of Tensor Flow computer vision models called Mobile Nets is focused on mobile devices and is intended to optimise accuracy while taking into account the limited resources of embedded or on-device applications. Mobile Nets are low-power, low-latency models that are sized and configured to satisfy different use-cases' resource requirements. They can serve as a foundation for segmentation, embeddings, detection, and classification

Experimental Result

The model's accuracy on the training data set is referred to as training accuracy. The model is trained on a set of labelled photos during the training phase. The accuracy is determined by the number of images the model properly classifies. When analyzing how successfully the model is classifying the photos in the training set, one helpful measure to have is the training accuracy. It's crucial to remember, nevertheless, that a high training accuracy does not guarantee that the model will function effectively when exposed to fresh, untested images. The model's accuracy on a different validation data set is known as validation accuracy. A section of the training data set is usually reserved as a validation set throughout the training phase.

Comparative Analysis of Image Classification Model

Key performance indicators for assessing the effectiveness of an image classification model are training and validation accuracy [21]. As the model learns from the training data, training accuracy quantifies the model's accuracy throughout this phase. During the validation phase, when the model's capacity to generalise to new data is examined, validation accuracy quantifies the model's accuracy on the unseen data. With an accuracy of 98.92% of the four transfer learning algorithms examined, Mobilenetv2 demonstrated the best level of accuracy. However, assessing the models' overall performance is difficult in the absence of training accuracy data. poor training accuracy could mean that the model needs more training data or changes, whereas high training accuracy and poor validation accuracy could mean that the model is overfitting the training set. In conclusion, while assessing an image classification model's performance, it is important to take into account both training and validation accuracy. Even if a high validation accuracy is preferred, it's crucial to make sure the training data isn't being over fitted by the model. To ascertain the model's total performance, it is crucial to assess both training and validation accuracy. Mobile net mobile net v2





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Accuracy

The metric quantifies the percentage of accurate classifications the model makes. Although it is a straightforward and simple statistic, it may be deceptive if the dataset is unbalanced, meaning that one type occurs substantially more frequently than the other.

Precision

The percentage of positive forecasts that come true is measured by this metric. This would refer to the percentage of leaf samples that are truly affected by the disease in the context of classifying grape leaf diseases. When the cost of false positives—that is, the prediction that a leaf has the illness when it does not—is high, precision becomes a valuable parameter.

CONCLUSION

When comparing the picture classification performance of the Mobile Net and MobileNetV2 models, MobileNetV2 has proven to perform better. Its superior computing power and efficiency overshadow Mobile Net, making it the go-to option for image classification applications. The extraordinary efficiency of MobileNetV2 is attributed to its architectural innovations, which include inverted residuals and efficient depth wise separable convolutions. These enable the network to deliver state-of-the-art results while spending fewer computational resources. This distinguishes MobileNetV2 as a unique model in the field of image categorization and demonstrates how well it strikes a balance between accuracy and efficiency. In order to effectively manage leaf diseases in agricultural crops, early detection and creative remedies are required. One potential solution to this issue is to integrate the use of frameworks such as MobileNetV2.Renowned for its remarkable precision and efficacy, MobileNetV2 has demonstrated its abilities in the field of image classification, rendering it a useful instrument for detecting leaf illnesses in farming crops and suggesting suitable remedies.

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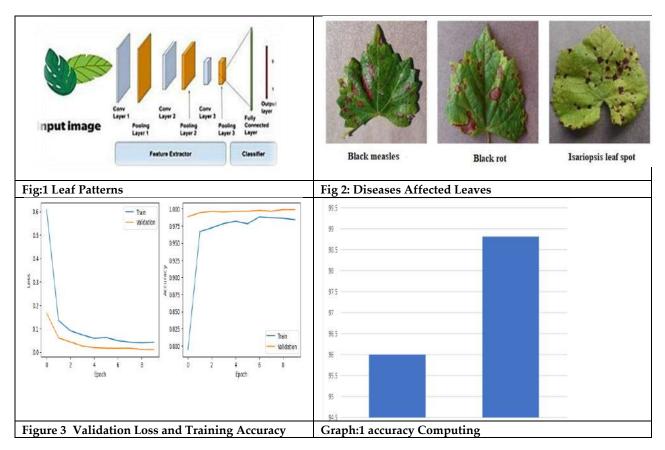


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Table1: Accuracy

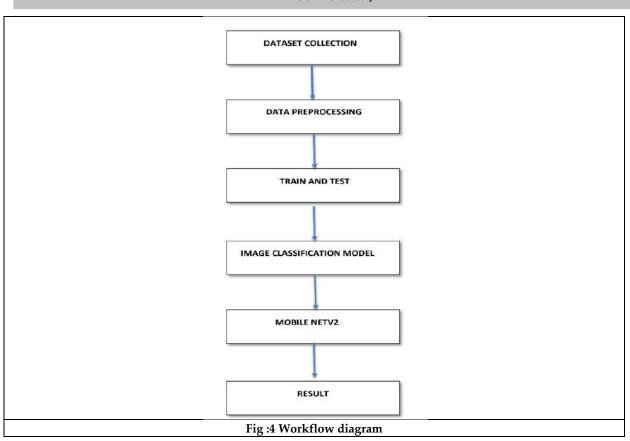
Model	Precision	Recall	F1 Score	
Mobile net	95.21	94.32	96.21	
Mobilenetv2	98.21	98.34	98.67	







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RESEARCH ARTICLE

Study of Image Segmentation of Color Image by Various Methods

P. Vinoth, P. Karthick* and T.Brinthaguru

Department of Mathematics, School of Advanced Science Kalasalingam Academy of Research and Education, Krishnankoil-626 126, Virudhunagar, Tamil Nadu, India.

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*Address for Correspondence

P. Karthick

Department of Mathematics, School of Advanced Science Kalasalingam Academy of Research and Education, Krishnankoil-626 126, Virudhunagar, Tamil Nadu, India. Email:karthickphd91@gmail.com



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ABSTRACT

Segmentation of images is critical for interpreting and evaluating items inside images. The procedure involves splitting and assessing images to convert them from vague to meaningful and helpful. This procedure is employed in a variety of disciplines, including medicine, culture, and industry. Image segmentation employs a wide range of functions, including edge and threshold functions. This dissertation will go through the color image segmentation strategies, present examples, and show the many sorts of images that may be used to find error ratio for better understand.

Keywords: Image segmentation, RGB image, Edge based method, Threshold method, Watershed method, Regionbased method, Clustering method.

INTRODUCTION

When it comes to the study and use of the picture, people are only interested in specific areas of it. These elements are sometimes referred to as the target or foreground (the other element is referred to as the background); they typically match the image in a particular and distinctive way. To identify and evaluate the item, it must extract and separate them; only then will it be able to be used for the target in the future. We have introduced the "image engineering" idea to demonstrate the degree of picture segmentation in image processing. This concept brings together the various theories, techniques, algorithms, tools, and equipment involved in image segmentation into acohesive whole. Image engineering is a relatively recent topic of study and application for images. Image segmentation is the process of dividing a digital image into several parts. The goal of segmentation is to separate an image in to more representative sections. Much simpler to examine. These areas could match certain surfaces, items, object' natural components. Typically, borders and objects are located through the process of picture segmentation. Based on the color characteristic of image pixels, color image segmentation presumes that objects with consistent colors in the





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image belong to distinct clusters and are therefore significant. Put otherwise, every cluster identifies a class of pixels that have comparable color characteristics.

RELATED WORK

Since there isn't a consensus on the optimum option for color space-based image segmentation, some study has attempted to determine which is the optimal color space for a particular assignment. There is no one color space that can produce segmentation results that are suitable for all types of photos since the segmentation results rely on the color space that is used. Because of this, numerous writers have attempted to identify the color space that will work best for their particular color image segmentation issue [1]. The color space, which enables the geometrical representation of colors in 2D and 3D space, is a mathematical representation of colors. Each pixel in an image can have its color information provided via color spaces. Color spaces provide helpful information on the appearance of the color spectrum, as seen in [2]. The fusion of color models and mapping functions is called color space. Each image has a set of intensity values, and noise might arise from arbitrary changes to these values. Typical noises include impulse noise, salt and pepper noise, and so on. Noise can lead to challenges in efficiently detecting edges; thus, image must be filtered to lower noise content that causes edge strength to be lost [3] Another name for its smoothing. Enhancement is the process of making an image better. Its goal is to create an image that is superior to the original and more appropriate. To improve the image's edge quality, a filter is used.

Here, morphologically-based image segmentation technique has been enhanced Under the complicated backdrop, image segmentation processing for wind turbine blades is done using a canny operator. Applying morphology to improve Canny operator's image segmentation method after Canny operator edge detection, the binary image is morphologically treated to eliminate redundant edge information, in contrast to the single image segmentation technique [4]. The phase of defining markers can be omitted during the watershed flooding change when a multiscale approach and region merging are used. We begin by computing the mean of each color class in the image to create an energy image. Deng introduces this criterion, which is known Automatic breast cancer diagnosis based on k-means clustering and adaptive thresholding hybrid segmentation [6]. It uses a texture discontinuity -based method to identify homogeneous areas in a picture. Every pixel in the complete image can have its measure computed. All colors are regarded as classes in the original Deng's version, and the color information is only utilized in a pre- processing quantization phase. The number of clusters must be known in advance for the k-means and fuzzy c- means algorithm and the user must provide the six parameters for the isolate algorithm [6].

Narayanamoorthy et all [11],[12] constructs, the new approach, which is based on the k-means algorithm, incorporates a validity metric based on the intra-cluster and inter-cluster distance measurements to get beyond the requirement of indicating the number of clusters. Furthermore, a slide prepared outside of normal protocol may result in under- or over-staining of the slide. Sharma et all, [7] Based on these justifications, the current study will make use of the color image segmentation approach's capability by utilizing a variety of color models and the kmeans clustering technique to produce the fully segmented image. Future study will concentrate on real-time segmentation and classification with the aid of more effective methods like ensemble learning and deep learning. Furthermore, we think that it will improve the efficacy and precision of algorithms used in object identification and image classification systems. In this paper [8] The collected results in dictated a noteworthy improvement in segmentation performance. The suggested approach may be helpful for color segmentation of images. In 2013 [9] represents, there are a few shortcomings with our suggested approach. The majority of the image models that are utilized are predicated on certain a priori information, such as the standard deviation and mean of each segmented image region. Furthermore, we have only taken into account a single image for every application in our work, even though the segmentation process may benefit greatly from many realizations of the same image fused together. This is because the validity score is significantly impacted by the much larger inter-cl user distance. [10] To get around this, first locate the validity measure's local maximum. Next, locate the value that is the minimum after the local maximum. The least number of clusters that can be chosen using this modified rule is four. This isn't really an issue because pictures in natural color.





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Proposed Methodology of Color Image Segmentation

In digital image processing and analysis, image segmentation is a commonly used technique for dividing animage into many parts or sections, usually based on the characteristics of the individual pixels in the image categories for segmenting images. There are various techniques for segmenting photos, including

- 1. Threshold Approach
- 2. Method based on region
- 3. Method based on edges
- 4. The watershed-based approach
- 5. A approach based on clustering

The RGB color space, which is the most often used color space, is defined by the three color components of the corresponding pixel, which are blue (B), green (G), and red (R). The color spaces were divided into the following groups. The real RGB, subtractive CMY, and imaginary XYZ primary spaces are the main spaces that are based on the theory that suggests it is feasible to match any color by mixing an approximate proportion of the three primary colors. The process of changing RGB to CMY is

C'=1-R	C = min(1, max(0, C' - K'))
M'=1-G	M = min(1, max(0, M' - K'))
Y' = 1 - B4	Y = min(1, max(0, Y' - K'))

$$K' = min(C', M', Y')$$

Segmentation based on threshold

Using this method, original photos can be converted to binary images. Pixels are compared to a preset threshold value in order to achieve this. pixels less than or equal to this value will be allocated 0 (white); pixels larger than this value will be assigned 1 (black). The following are the steps in the algorithm:

- 1. Read an image as input.
- 2. Make new variables in two dimensions and assign the picture size to them.
- 3. Type in the threshold amount.
- 4. Construct a fresh zeros matrix that has the same measurements as the picture.
- 5. Align the pixels with the threshold amount.
- 6. Display the outcomes.

We must jot down the elements in the new image if we need to split them up.

- Step 1: Use an input image
- Step 2: Make this picture grayscale. Step
- Step 3: Give the value for the threshold.
- Step 4: Only display pixels that are larger than the threshold.

Segmentation based on edges

This technique uses intensity-level (grayscale) detection to extract object information. One of the methods most frequently used to analyse photos is edge detection. The edge method determines the features of the edges by comparing colors. These details are displayed as black backgrounds with white lines. There are numerous approaches for filtering

- Sobel filters
- Prewitt filter
- · Robert's filter
- Log filter





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· zero-cross screening.

The variable in the image and the procedure are the two parameters of this function (the Edge function).

The algorithm's steps are as follows:

- 1. Examine a picture.
- 2. Apply any of the aforementioned filters along with the edge function.
- 3. Make the image visible

Segmentation of watersheds

When splitting items that are in contact with one another, watersheds offer a free object-splitting method that is quite helpful. Images are interpreted by this technique as physiological surfaces, whose elevation is represented by the value of f (x, y). The watershed algorithm locates the hill lines and watershed basins in the picture where water would presumably accumulate in theory. Assume that each regional low point has a perforated hole, and that water rises through these holes at a constant rate, submerging the natural terrain below. As the water level rises, pixels are designated as submerged. Water will the entire image gets split into distinct collection basins as the flood goes on.

This algorithm's steps are as follows:

Step1: Read an image.

Step 2: Remove the black border before processing.

Step3: Modify the region within the watershed.

Step4:Calculate the Lesion Ratio.

Step5: Combine watershed regions.

Step 6: Define the boundaries of watersheds.

Step 7: Smooth borders (b-spline smoothing and border sampling).

Step 8: Add a border overlay

Segmentation based on clustering

In data mining, clustering is an undirected technique that helps find multiple latent patterns in the data without generating a specific hypothesis. The goal of clustering is to find commonalities amongst individual objects and create a collection of related ones. Hierarchical and non-hierarchical clustering techniques are the two forms of clustering. Non-classical Grouping This approach divides the dataset into M clusters, each having N items. K-means is the most used non-hierarchical clustering method in business intelligence. Grouping in Hierarchies This process results in a collection of nested clusters. Every pair of objects in these nested clusters is further nested to create alarger cluster until, at the end, only one cluster is left. K stands for Clustering Technique. K-means clustering is an unsupervised method that uses the input data without a labeled response. One popular method for clustering is K-means clustering. Typically, practitioners first, familiarize yourself with the architecture of the dataset. K-means is used to cluster data points into discrete, non-overlapping groups.

RESULTS AND DISCUSSIONS

A set of photos and a few commonly used algorithms from the literature are used to gauge how well the recommended algorithm is implemented. Real photos are utilized for performance estimation and comparisons, when the item can be precisely separated from the background using a suitable threshold technique. The illustrations can be found in figures 1–5. Each figure displayed the following: (a) the original image; (b) the original image's histogram; (c) the threshold image; (d) the edge-based image; (f) the watershed image; and (e) the K mean. clustering technique for segmenting images.

Ratio of Error (ER)

The most important comparative indicator is the error ratio (ER). ER is the ratio of pixels in the erroneous area





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between the object pixels and the optimal threshold image produced by each approach. incorrect segmentation as indicated by the formula below,

$$E_{R=\frac{N_r+}{N_r}\times 100}$$

where NR is the number of pixels in the segmented pictures and Nm is and Nr are the number of segmentation pixels in the erroneous area. In this sense, the ER value is a number between 0 and 100; a value of 0 indicates that there is no similarity between the segmentation result and the ground truth image. A excellent segmentation result is likewise indicated by a value of 100. The error shows the numerical numbers that, in each instance, correspond to how comparable the edge-based approach and the watershed method (k means clustering method) are. Next, we select the best error ratio number from a range of values. The lowest error value has been seen on bold type. Note that in each and everyinstance.

CONCLUSION

Image segmentation algorithms are able to convert indistinct photos into ones that are interpretable enough to extract important information, especially in the medical field. This study looked at the performance and error rate of a few MATLAB-based hashing algorithms. It is important to note that not all of the photos we utilized haditems detected by these techniques. On the other hand, low-density pixels or photos with poor color density were simply analysed. An efficient evaluation of uniformity was not possible. To find the most efficient picture segmentation system, researchers must keep applying deep learning techniques.

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Table 1: Error Ratio

Images	Threshold Method		Error Ratio (with%)			
	Original Image	Optimum Image	Edge Based	K means Clustered	Watershed	
	pixel	Pixel	Method	Method	Method	
Fox.jpg	1,998,000	666,000	28.8%	25.3%	10.2%	
Parrot.jpg	2,281,248	760,416	13.65%	8.9%	9.2%	
Sunflower.jpg	817,920	272,640	8.1%	6.5%	18.5%	



Figure 1: Image Segmentation Comparison for Fox Image (a) Original Image

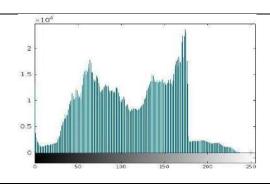


Figure 1: (b) Histogram of the Original Image



Figure 1: (c) Threshold Image



Figure 1: (d) Edge based Image





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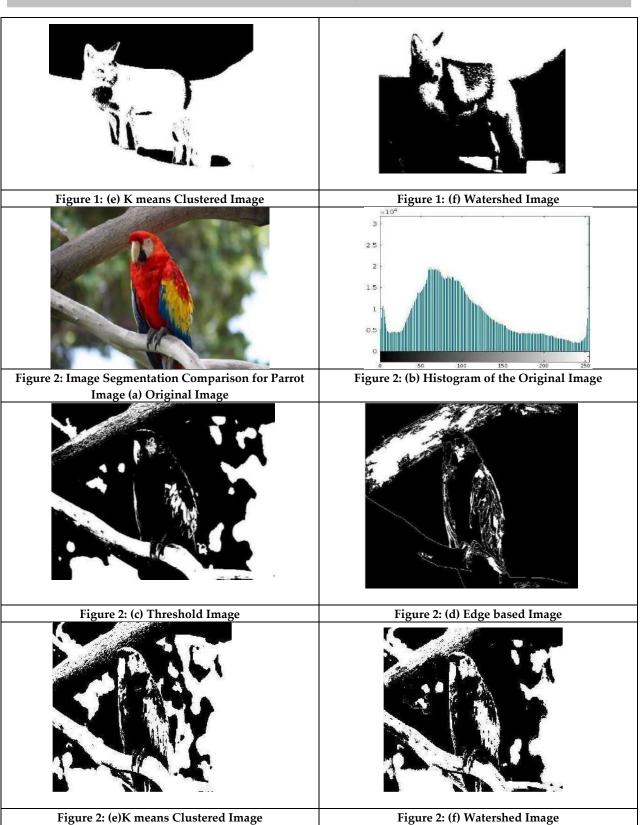




Figure 3: (e) K means Watershed Image



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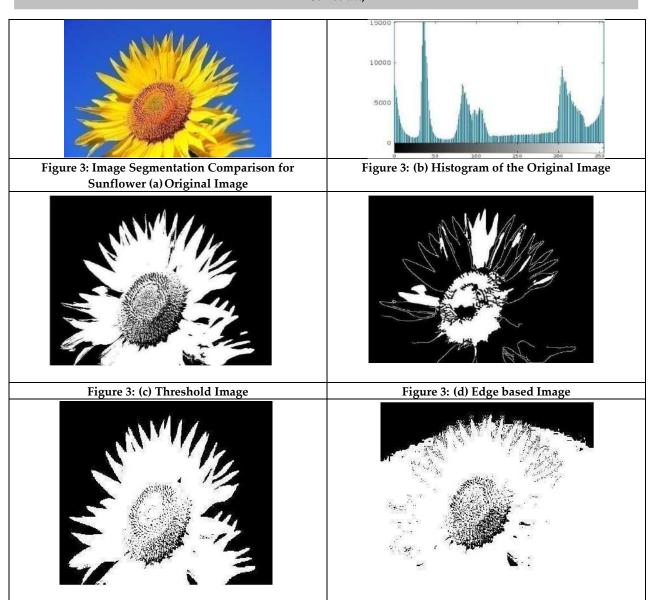
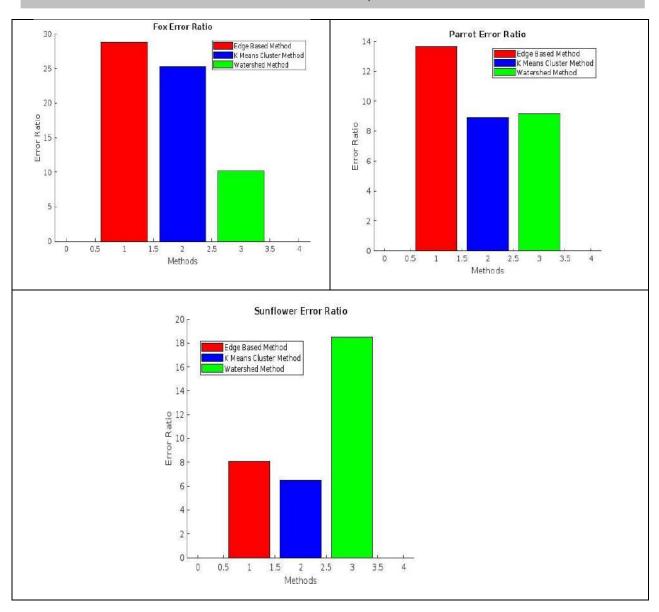




Figure 3: (f) Watershed Image



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RESEARCH ARTICLE

Memory Mate: A Mobile Application for Empowering Alzheimer's Patients

M.Sridhar^{1*} and R.Sankaran²

¹Assistant Professor, Department of Information Technology, Sri Ramakrishna College of Arts & Science, Coimbatore, Tamil Nadu, India.

²PG Student, Department of Information Technology, Sri Ramakrishna College of Arts & Science, Coimbatore, Tamil Nadu, India.

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*Address for Correspondence

M.Sridhar

Assistant Professor, Department of Information Technology, Sri Ramakrishna College of Arts & Science, Coimbatore, Tamil Nadu, India. Email: sridhar@srcas.ac.in



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ABSTRACT

A constant clinical decline over several years is an indication of Alzheimer's disease. It impairs daily task performance and induces memory loss. Memory problems can make it difficult to recall names, faces, locations, or other details of people. Thus, the purpose of our study is to support the continued independence and social participation of individuals with moderate (the initial stages) and intermediate (middle-stage) Alzheimer's disease. We propose an Android app that uses facial recognition and position detection from Google Maps. The paper aims to improve users' ability to perform daily tasks and to enhance ordinary communication by integrating a notification element. By tracking their movements and prevents them from becoming lost, the Global Positioning System (GPS) detection feature helps keep dementia patients safe. The application has consistently assisted persons with signs of Alzheimer's and significantly enhanced their quality of life, according to the results. Therefore, our findings highlights the need of utilising artificial intelligence (AI)-based features—such as facial recognition in this case—when developing healthcare applications that could significantly impact society.

Keywords: Face recognition, Alzheimer's illness, smartphone applications and Machine learning

INTRODUCTION

The fact that aging is a recognized risk factor for Alzheimer's disease and that more people will likely develop the illness as the population ages, the disease is becoming a growing public health problem. It primarily affects the elderly, some experts refer to it as the "century disease." Simple daily activities like eating and cleaning their teeth





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might be difficult for those who have Alzheimer's. Alzheimer's disease is a severe, long-term brain illness that starts with memory loss and progresses to planning, thinking, language, concentration, and cognition problems. Therefore, the goal of this application was to help families and patients with Alzheimer's disease live better lives by developing a smartphone application that uses face recognition technology to enable machine learning. Nowadays, face recognition is being used in many real-world applications, from personal to public security. It is employed to confirm someone's identify or to locate and follow a potential suspect in a crowd. Globally, Alzheimer's disease is becoming a more significant problem for individuals, their households, and healthcare systems due to its relentless and debilitating nature. Alzheimer's disease is becoming more common as the world's population ages, necessitating the development of creative solutions to improve the condition of those who are afflicted and assist those who are caring for them. In the current digital era, smartphone apps have become extremely effective resources for managing the particular requirements and difficulties related to Alzheimer's care. This study uses the features of the Android framework to construct a care application specifically for patients with Alzheimer's disease. This causes people to lose their independence, memory, and cognitive function, which makes it harder to carry out daily chores and activities. With compassion and knowledge, the suggested application aims to close this gap by offering a feature-rich package catered to the unique requirements of Alzheimer's sufferers and their carer givers. Because of its adaptability and cross-platform interoperability, Android is a great platform for developing applications that are simple to use and intuitive for users. The application's seamless functioning on both iOS and Android devices guarantees its universal accessibility and makes it easier for individuals with Alzheimer's disease and those who provide care for them to use. This introduction chapter establishes the background against which we will examine the development of this essential care application. Examine the unique challenges faced by dementia patients and their carer givers, with a focus on how mobile technology might be able to alleviate some of these challenges. The study's main objective is to assist individuals with Alzheimer's disease who are in the early or middle stages of the illness by providing them with a tracking system, daily task reminders, social engagement opportunities, and increased confidence in their ability to remember each member of their family. The proposed application could help all mild-to-moderate Alzheimer's sufferers and their careers by developing an application with features like machine learning-driven facial recognition. This helps the patient's close friends and family remember things and provides them with a smart GPS tracking device that can be used to help carers find the patient. The intended use aims to improve the standards of life for individuals with Alzheimer's disease by simplifying their lives and people around them.

LITERATURE SURVEY

Widely used to describe a range of conditions, the term "dementia" is most frequently linked to impairments in language, recall, cognition, and problem-solving abilities. One's capacity to carry out regular tasks is thereby compromised. Alzheimer's disease is the most common form of dementia, and its incidence has been rising in recent years. Research is increasingly demonstrating that advancements in technology have a major impact on individuals with Alzheimer's disease, especially during the early stages of the illness. Algorithms and intelligent devices can help ease some of the challenges dementia patients and caretakers frequently encounter when providing supportive care. Today, the GPS is a widely used global navigation device. It was initially developed by the US United States Department of Defence (USDOD) utilising signals from satellites. The system consists of a minimum of 24 satellites. Since there are no installation or subscription costs, it is a very versatile and reasonably priced instrument that can be used in any weather and anywhere in the world. The satellites were first put into orbit by the USDOD solely for military purposes, but in the 1980s, these were extended to include public coverage as well. GPS allows for the precise location of latitude and longitude coordinates at ground level. The method is calculating and determining the current location by examining the differences in timing between signals coming from thirty satellites that are under the management of USDOD and numerous commercial organisations. It has also been demonstrated to be a successful method for resolving problems with outdoor localization because GPS signals typically demand an unbroken receiversatellite line-of-sight (LOS) channel. Automatic facial recognition is one of the key advancements impacting biometrics technology, artificial intelligence, and machine learning. Their effect has mostly been attributed to their broad range of potential applications and the current focus on them by the scientific community. In essence, facial recognition devices





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are two pieces of technology that do two main jobs: they identify the individual whose picture or video is used to confirm their identity, and they also identify the individual. In recent years, they have been employed as a means of controlling access in security systems. Because of this and the variety of uses they have, these systems have become the focus of numerous studies. The idea of machine learning was largely inspired by research into the interactions and communication that occurs between brain nerve cells through excitable neural activity. Since then, the popularity of machine learning models has grown, in part due to their problem-solving abilities. In more recent times, they have contributed to a number of very important technological developments. One of the primary objectives in the field of machine learning is to create computer systems that can learn from experience and develop on their own. As a result, AI machine learning has emerged as the preferred option for software developers operating in numerous domains, such as computer vision, facial recognition, natural language processing, robotics control, and numerous others. Consequently, machine learning is used to address each of the crucial phases of the recognition of faces process. This entails considering a variety of elements, such as the fact that faces are not just a collection of pixel-by-pixel images but rather have a distinct structure and symmetry. These systems can analyse sample facial photos that are kept in databases, capturing their distinct pattern features and cross-referencing them with particular images that are kept. The Facial Identification Grand Challenge was first introduced in 2006 to determine whether face recognition systems based on models developed using machine learning were accomplishing their stated goals.

PROPOSED SYSTEM

The technique in this paper was developed specifically to achieve its objective. Thus, the phases that are involved include data collection, database design, application interface design, implementation, and testing. For the purpose of this study, a web-based questionnaire was designed and disseminated to gather information from the main demographic targeted for this application-those who provide care for people with Alzheimer's disease. The questionnaire was designed to collect data regarding technology support services available to relatives and individuals suffering from Alzheimer's disease. The survey was created using Google Forms. The functional requirements list the tasks that a system has to be able to perform. They essentially outline the proper way for a system to function in specific scenarios. The application should display a list of related photos along with background information regarding each image's relationship to the patient. The patient should receive alerts and reminders from the app about important tasks that need to be finished. The application should notify the patient's carer whenever an indication from the surveillance bracelet shows that the individual has outside a designated safe zone. Non-functional requirements specify the performance limitations and the criteria using which the system will be assessed. The nonfunctional needs of performance and usability are crucial. The architecture shows how the needs and the created system align. Two separate architectural portions serve both the client and the care giver. The application has user interface components for adding, editing, and comparing pictures from the patient's perspective. The pictures are kept inside a central database that is accessible by other sections. Carers can add photos, compare photos, specify and monitor the location, and set reminders with this application. The design step streamlines the process by taking the requirements from the evaluation phase and putting them into an architectural chart. This graphical representation includes information on the necessary components and how they interact. Furthermore, this phase solves the problem of how to design the best potential answer with the aid of prototype development. Therefore, the primary goal of the designing phase is to facilitate the development of an interface that allows the systems to function as intended across the board. When a fresh model or a modified version of a previous one is being developed, an early sample or early model is called a working model in the realm of software development.

RESULTS AND DISCUSSION

The Memory Mate program's results show that it can identify people by looking through an album that has categorised each person by face and displayed information about their relationship. When it comes to ensuring that users can take full advantage of an application's capabilities and make money from it, its usability is paramount. An exploratory survey with one hundred users helped determine how useful this programme is. Users were required to





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utilise the application before being given a series of questions to respond to in order to get input on different usability-related issues. The questionnaire items that were utilised to evaluate the usefulness were based on the McLaughlin and Skinner principles. The process of usability testing produced the following results:

- 1. Alzheimer's Assistant has a validation system in place to ensure that data is accurate.
- 2. Users are confident in their capacity to use it and believe it to be effective.
- 3. Users claim to be able to operate the system, particularly when supplying or extracting data.
- 4. The application is easy to use; it is believed to be a quick and effective system.
- 5. The information it produces is easy to understand.

CONCLUSION

An app called Alzheimer Assistant was created especially to support those who have Alzheimer's disease. It can send patients notifications to remind them of daily tasks; this can help detect individuals by relating to a record in which every individual has been categorised through face and his connection information displayed; and it can give patients a GPS bracelet to ensure their family can find them precisely in the event that they get lost. Consequently, this programme helps people with Alzheimer's disease feel more confident when going about their everyday lives, which improves their ability to routinely go to social events. The ability of this application to assist families and care givers has been extensively demonstrated. The application's primary flaw prevents carers from remotely monitoring patients, which prevents the Alzheimer's Patients Association from formally endorsing it. The usefulness of the application would therefore be improved by creating and implementing this new feature, which would be included in later releases. Future research projects will focus on deep learning algorithms for facial recognition because of their significant impact in this field and their capacity to provide outcomes with superhuman performance and accuracy. The suite of software tools previously available to assist individuals with Alzheimer's disease in preserving social connections with their loved ones is enhanced by the Alzheimer Assistant app.

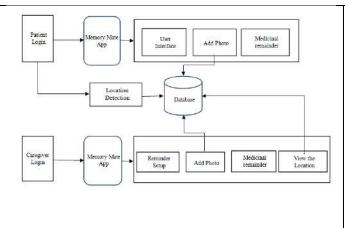
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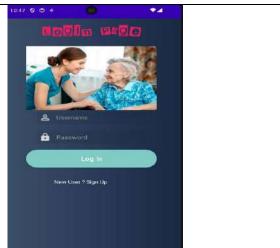


Figure 1: Architecture Design of the Application



Figure 2: Login page



Figure 3: Registration page





Figure 5: Medicine page and Location tracking





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REVIEW ARTICLE

Detection and Classification of Leaves in a Video: A Literature Review

Vidyashankara^{1*}, Hemantha Kumar G¹, Navena M³, Sumanashree Y S² and Mallesh R ³

- ¹Department of Studies in Computer Science, University of Mysore, Mysuru, Karnataka, India.
- ²PG Department of Computer Science, JSS College (Autonomous), Mysuru, Karnataka, India.
- ³Department of Computer Science, JSS College for Women's, Chamararaja Nagara, Karnataka, India.

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*Address for Correspondence Vidyashankara

Department of Studies in Computer Science, University of Mysore, Mysuru, Karnataka, India. Email: vidhyashankara.s@gmail.com



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ABSTRACT

In recent years, the computer vision field has made significant advancements in various domains, including video analysis. One particular area of interest is leaf detection and classification, which plays a crucial role in agriculture, environmental monitoring, and plant disease diagnosis. Being able to automatically analyze leaves from video data has significant applications across agriculture, botany, ecology, and other domains. Specifically, in agriculture, detecting and classifying leaves through computer vision techniques enables non-invasive monitoring of crop growth and plant health at scale. Leaf analysis is challenging because of factors like cluttered backgrounds, leaf occlusion, and lighting changes. Early works focused on analyzing individual static images of leaves, but recent advances in computer vision and deep learning have enabled the application of leaf analysis techniques to video streams. Analyzing leaf data in the video provides benefits like capturing temporal changes, wider observational coverage, and 3D structure clues through the direction of frame motion. This paper presents a comprehensive review of the recent advancements in the field of detection and classification of leaves in a video. The paper discusses the various techniques used for detecting and classifying leaves in images and the need for video Analysis & sampling, including image processing, machine learning, and deep learning. The paper also highlights the challenges faced in this field and the future scope of research.

Keywords: cluttered backgrounds, leaf occlusion, temporal, 3D structure, image processing, machine learning, deep learning, video analysis





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INTRODUCTION

Plants fulfill a significant role in human existence through their provision of shelter and facilitation of a healthy and breathable environment. The vast number of plant species found across the globe necessitates the creation of a comprehensive plant database, enabling swift and efficient classification and recognition methods. The environment and climate are heavily reliant on the presence of plants [25], and individuals involved in natural resource management, particularly those engaged in wildlife conservation, must possess the ability to determine whether a variety of plant species are present or absent. In light of the limited knowledge regarding plants [17, 26], an innovative plant recognition system has been developed using digital images and videos of plant leaves. Given the increasing scarcity of plant categories and the impending extinction of many plant species [12], there is a pressing need for a simple and effective means of recognizing and classifying plants based on their respective categories. It is also helpful for managing and maintaining plants through the early detection of plant diseases.

Leaf detection and classification are important tasks in a variety of applications, such as:

Plant phenotyping

Leaf detection and classification can be used to measure plant growth and development, as well as to identify plant diseases [29]. This information can be used to improve crop yields and develop new disease-resistant varieties.

Precision agriculture

Leaf detection and classification can identify pests and diseases early on [35, 36]. This information can be used to target pesticides and fertilizers more precisely, thereby reducing costs and environmental impact [3].

Environmental monitoring

Leaf detection and classification can be used to monitor the health of forests and other ecosystems [2, 34]. This information can be used to track the spread of invasive plant species and to assess the impact of climate change. Accurate and efficient leaf detection and classification are essential for enabling these applications.

Identification of specific plant types relies on discerning characteristics such as Stem, flower, fruit, and leaf characteristics like leaf shape, colour, and texture [1, 14]. Major research activities are based on leaf characteristics, because of availability in large quantities and easy sample collection and these activities do not adversely affect existing plants and the ecosystem.

Recognition System

A recognition system is a type of artificial intelligence (AI) system that can identify and classify objects or patterns. This system is shown in Fig. 1. It typically follows a Four-stage process:

Data Acquisition

The first step involves acquiring the input data. This data can be in various forms, such as images, audio, or video. The quality and relevance of the input data significantly impact the performance of the recognition system.

Preprocessing

The image is Preprocessed to remove noise and enhance the features of the leaves. This may include steps such as contrast enhancement, histogram equalization, and filtering. It also covers segmentation. Where the image is segmented to isolate the leaves from the background. This is typically done using techniques such as thresholding, edge detection, and region growing.





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Feature Extraction

The second step involves extracting relevant features from the acquired data. These features represent the essential characteristics of the input data that can be used for identification or classification. The choice of features depends on the specific recognition task and the nature of the input data.

Pattern Matching or Classification

The third step involves matching the extracted features to a database of known patterns or classifying the input data into predefined categories. This involves comparing the features to reference patterns or using machine learning algorithms to make predictions.

Deep Learning-Based Leaf Detection Methods

Deep learning-based leaf detection methods have recently emerged as a promising approach. These methods utilize deep neural networks to learn features from data without the need for manual feature engineering. Deep learning-based leaf detection methods have achieved state-of-the-art results on both image and video datasets. Leaf recognition using a deep learning approach typically involves the following steps:

Data Collection and Pre-processing

Gather a diverse dataset of leaf images

Collect a large and diverse dataset of leaf images representing different species, shapes, sizes, and conditions. Ensure the dataset includes images with varying illumination, backgrounds, and occlusions to enhance the model's robustness.

Pre-process the images

Pre-process the leaf images to ensure consistency in size, format, and color balance. This may involve resizing, normalizing pixel values, and adjusting contrast and brightness.

Data augmentation

Augment the dataset to increase its variability and improve model generalization. Apply techniques like flipping, rotating, cropping, and random distortions to create more variations of the existing images.

Model Architecture Selection

Choose a deep learning architecture

Select a suitable deep learning architecture for image classification, such as convolutional neural networks (CNNs). CNNs are particularly effective in extracting features from images and learning complex patterns.

Consider transfer learning

Utilize transfer learning by leveraging pre-trained CNN models like VGG16, ResNet, or Inception. These models have already learned generic features from large image datasets and can be fine-tuned for leaf recognition.

Model Training

Split the dataset

Divide the pre-processed dataset into training, validation, and testing sets. The training set is used to train the model, the validation set is used to monitor overfitting and tune hyperparameters, and the testing set is used to evaluate the final model's performance.

Define the loss function

Choose an appropriate loss function for image classification, such as categorical cross-entropy loss. The loss function measures the difference between the model's predictions and the true labels.

Optimize the model

Use an optimization algorithm like stochastic gradient descent (SGD) or Adam to update the model's parameters and minimize the loss function.

Hyperparameter tuning

Tune hyperparameters like learning rate, batch size, and optimizer settings to optimize the model's performance on the validation set.





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Model Evaluation and Testing

Evaluate the model on the testing set

Assess the model's generalization ability and real-world performance by evaluating its accuracy, precision, recall, and F1 score on the testing set.

Analyze errors

Identify and analyze the types of errors the model makes to understand its limitations and areas for improvement.

Deployment and Application

Integrate the model

Integrate the trained deep learning model into a practical application, such as a leaf identification tool or an autonomous agricultural robot.

Continuous improvement

Continuously monitor the model's performance in real world scenarios and collect new data to retrain and improve the model as needed.

Deep learning offers several advantages for leaf recognition

Automatic Feature Learning

CNNs can automatically extract and learn relevant features from leaf images without the need for manual feature engineering.

Robustness to Variations

CNNs are relatively robust to variations in leaf appearance, such as lighting conditions, background clutter, and partial occlusions.

High Accuracy

Deep learning models have achieved state-of-the-art accuracy in leaf recognition tasks. The deep learning approach has revolutionized leaf recognition, enabling more accurate and efficient leaf identification in a wide range of applications.

The Leaf detection and classification typically follow a two-stages:

Leaf detection

A Method is used to detect leaves in the video frames or image and extract them [38]. This method is useful for identifying required objects of interest in a frame and extracting them by removing unwanted complex backgrounds. There are several leaf detection algorithms available Here are some of them:

Feature detection algorithms

These algorithms are used to detect specific features in an image such as corners, blobs, circles, and so on. Some of the popular feature detection algorithms include Harris Corner Detection, Shi-Tomasi Corner Detector, Scale-Invariant Feature Transform (SIFT), and Speeded-up Robust Features (SURF) [44]. The advantages of feature detection algorithms are that they are computationally efficient and can be used for real time applications. However, they are sensitive to changes in lighting conditions and may not work well with complex images.

Object detection algorithms

These algorithms are used to detect objects in an image and classify them into different categories. Some of the popular object detection algorithms include Fast R-CNN, Faster R-CNN, Histogram of Oriented Gradients (HOG), Region-based Convolutional Neural Networks (R-CNN), Region-based Fully Convolutional Network (R-FCN), Single Shot Detector (SSD), Spatial Pyramid Pooling (SPP-net), and You Only Look Once (YOLO) [45]. The advantages of object detection algorithms are that they are highly accurate and can work well with complex images. However, they are computationally expensive and may not be suitable for real-time applications.





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Image segmentation algorithms

These algorithms are used to partition an image into multiple segments or regions based on the similarity of the pixels in the image. Some of the popular image segmentation algorithms include the Watershed Algorithm, Mean Shift Algorithm, and Normalized Cuts Algorithm [46]. The advantages of image segmentation algorithms are that they can be used to extract meaningful information from an image and can work well with complex images. However, they are computationally expensive and may not be suitable for real-time applications. some detection methods available for video samples:

Object tracking

This method involves tracking an object in a video stream or a video sequence by considering the current image and the previous ones.

Action classification

This method involves classifying the actions performed by an object in a video sequence.

Optical flow estimation

This method involves computing the motion of objects in a video sequence by computing the pixel shift between two frames.

Trajectory classification

This method involves classifying the trajectories of objects in a video sequence [48].

Low rank sparse matrix

This method involves decomposing a video sequence into a low-rank matrix and a sparse matrix to detect objects [48].

Background subtraction

This method involves subtracting the background from a video sequence to detect moving objects [48].

Object detection using deep learning

This method involves using deep learning algorithms such as Convolutional Neural Networks (CNN) to detect objects in images and videos [49].

Leaf classification

Once the leaves are detected, this method is useful for classifying them based on their extracted features [39, 42]. This method classifies leaves by typically the percentage of matching the features extracted from the samples at the time of testing with the features trained on a dataset of labelled leaf images from different species. There are several leaf classification algorithms. Here are some of them: The Random Forest algorithm is a technique for ensemble learning, wherein multiple decision trees are created during the training phase. The resulting output is determined by the class that appears most frequently among the individual trees in the case of classification, or by the average prediction in the case of regression [53]. The advantages of Random Forest are that it is highly accurate, can handle missing data, and can be used for feature selection. However, the execution of this task incurs a significant computational cost and may exhibit incompatibility with time-sensitive applications.

Support Vector Machines (SVM)

This algorithm is a classifier, either linear or non-linear in nature, which is determined by the kernel employed. In the case of utilizing a linear kernel, the classifier and consequently the prediction boundary exhibit linearity. In order to distinguish between two classes, it is imperative to establish a line that possesses a maximum margin. This particular line is drawn equidistant from both sets. Additionally, two supplementary lines are drawn on each side, referred to as support vectors. SVMs learn from the support vectors, unlike other machine learning models that learn from the correct and incorrect data [52]. The strengths of Support Vector Machine (SVM) lie in its notable precision, ability to process data with a high number of dimensions, and efficacy in handling limited datasets. Nevertheless, it is crucial to acknowledge that SVM incurs substantial computational costs and may not be optimal for applications requiring real-time responsiveness.





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K-Nearest Neighbors (KNN)

This algorithm is a nonlinear classifier that predicts which class a new test data point belongs to by identifying its k nearest neighbors class. We choose these k closest neighbors by considering the Euclidean distance metric. Within this group of k neighbors, we tally the quantity of data points belonging to each category, and subsequently assign the new data point to the category that has the highest number of neighboring points [52]. The advantages of KNN are that it is simple, easy to implement, and can work well with small datasets. However, it is computationally expensive and may not be suitable for high dimensional data.

Convolutional Neural Networks (CNN)

This algorithm is a deep learning architecture that is used for image classification. CNNs are designed to automatically and adaptively learn spatial hierarchies of features from input images by enforcing a local connectivity pattern between neurons of adjacent layers [52]. The advantages of CNN are that it is highly accurate, can handle complex images, and can work well with large datasets. However, it is computationally expensive and requires a large amount of training data.

Recurrent Neural Networks (RNN)

This algorithm is a deep learning architecture that is used for sequential data such as videos. RNNs are designed to process sequential data by maintaining an internal state or memory of the previous inputs [54]. The advantages of RNN are that it can handle sequential data, can work well with variable-length inputs, and can be used for real-time applications. However, it is computationally expensive and may suffer from the vanishing gradient problem.

Long Short-Term Memory (LSTM)

This algorithm is a type of RNN that is designed to overcome the vanishing gradient problem. LSTMs are designed to maintain an internal state or memory of the previous inputs and selectively forget or remember certain inputs based on their importance [51]. The advantages of LSTM are that it can handle sequential data, can work well with variable-length inputs, and can be used for real-time applications. However, it is computationally expensive and may require a large amount of training data.

Two-Stream Convolutional Neural Networks

This algorithm is a deep learning architecture that is used for video classification. It consists of two separate CNNs, one for spatial information and one for temporal information [53]. The advantages of Two-Stream CNNs are that it can handle complex videos, can work well with large datasets, and can be used for real-time applications. However, it is computationally expensive and requires a large amount of training data. However, leaf detection and classification can be challenging due to the complex and varying nature of leaf appearances [37]. Leaves can exhibit a wide range of shapes, sizes, colors, and textures. The plant leaves have high intra class variability and sometimes the leaves of different plants are very similar, which makes this task difficult even for botanists [16]. In addition, leaves can be occluded by other objects, and their appearance can change due to varying illumination conditions. Existing methods for leaf detection and classification have achieved promising results on leaf images. However, these methods often rely on hand-crafted features that may not be robust to the aforementioned challenges. In addition, existing methods are often computationally expensive, limiting their applicability to real time applications [33].

Related Work

There have been significant research approaches carried out in the first few years of the last two decades on the detection and classification of leaves, but later on, they concentrated on plant diseases or leaf disease recognition. Computer-aided plant recognition is still a very challenging task in computer vision due to the lack of proper models or representation schemes [20]. The objective of computerized identification of live plants was to quantify the features based on leaf geometry, morphology, and Fourier moments [55] before the introduction of Machine Learning and Deep Learning. These works can be mainly categorized as follows:





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Approaches for leaf detection and classification

Traditional Image Processing Techniques: These techniques involve the use of filters, thresholding, and morphological operations to extract features from leaf images. The extracted features are then used to classify the leaves into different categories [5,29]. Image Processing and Pattern Recognition techniques such as segmentation, feature extraction, and classification are commonly used for detecting and classifying leaves. Many scholars have endeavored to create a more resilient and effective system for identifying plants by utilizing techniques in pattern recognition and image processing that are centered around plant leaves, flowers, barks, and fruits [23]. Nevertheless, leaves hold a paramount significance compared to other plant components due to their abundance of valuable information and heightened dependability. Machine Learning-based approaches: These approaches use machine learning algorithms to learn the features of the leaf images and classify them into different categories. Some of the popular machine learning algorithms used for leaf classification include Random Forest, Support Vector Machines (SVM), and Convolutional Neural Networks (CNN) [27, 31].

Deep Learning-based approaches

These approaches use deep neural networks to learn the features of the leaf images and classify them into different categories. Some of the popular deep learning architectures used for leaf classification include AlexNet, VGGNet, and ResNet

Hybrid approaches

These approaches combine traditional image processing techniques with machine

learning or deep learning algorithms to improve the accuracy of leaf classification [20, 32]. As shown in fig. No. 2. A variety of methods have been proposed for leaf detection and classification in images [43]. Early traditional methods relied on hand-crafted features, such as color, shape, and texture, to represent leaves [42, 23]. However, these features are not always robust to the challenges mentioned above. In recent years, machine learning techniques such as support vector machines (SVM), decision trees, and random forests have also been used for detecting and classifying leaves. Deep learning techniques such as convolutional neural networks (CNN) have shown promising results for detecting and classifying leaves [46, 39]. Deep learning-based methods have achieved state-of-the-art results in leaf detection and classification. Deep learning methods can learn complex features from data without the need for hand-crafted features [31, 32, 40]. This makes deep learning methods more robust to the challenges of leaf detection and classification.

Deep Learning Methods

Several deep learning-based methods have been developed for leaf detection and classification [40]. These methods can be broadly divided into two categories: Two-stage methods: Two-stage methods first detect leaves in a video frame and then classify the detected leaves. A common example of a two-stage method is the Faster Region-based Convolutional Neural Network (Faster R-CNN) [41].

Single-stage methods

Single-stage methods detect and classify leaves in a video frame in a single step. A common example of a single-stage method is the You Only Look Once (YOLO) algorithm [45]. One of the most popular deep learning-based methods for leaf detection is the Faster Region-based Convolutional Neural Network (Faster R-CNN). Faster R-CNN is a two-stage object detection method that first generates a set of candidate's bounding boxes and then classifies each bounding box. Faster R-CNN has been shown to achieve good performance on leaf detection in videos. Another popular deep learning-based method for leaf detection is the Single Shot Multi-Box Detector (SSD). SSD is a one-stage object detection method that directly predicts bounding boxes and class labels for each pixel in an image. SSD is faster than Faster R-CNN, but it may not be as accurate for leaf detection in videos. such as the Convolutional Neural Network (CNN) and the Support Vector Machine (SVM). CNNs have been shown to achieve good performance on leaf classification tasks, especially when trained on large datasets of leaf images. SVMs are also effective for leaf classification, but they may not be as accurate as CNNs when trained on small datasets. Deep learning-based methods have achieved state-of-the art results on leaf detection and classification benchmarks [41].





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For example, the YOLOv5 algorithm achieved an accuracy of over 99% on the public LeafNet dataset. Recurrent neural networks (RNNs) are a powerful model for sequential data. End-to-end training methods such as Connectionist Temporal Classification make it possible to train RNNs for sequence labeling problems where the input-output alignment is unknown [32]. However, there are still many challenges that need to be addressed in the field of leaf detection and classification [35].

Proposed Methodology and Discussion

Video analysis is a rapidly evolving field that uses computer algorithms and techniques to extract meaningful information from video data. It has gained significant attention in recent years due to its wide range of applications across various domains, including surveillance, sports analytics, healthcare, and more. Video analysis algorithms can be used to detect and track objects, recognize faces, and identify suspicious activities in real-time, making security systems more proactive and efficient [33]. Leaf detection and classification in video analysis play a crucial role in various fields, such as agriculture, environmental monitoring, and plant disease detection [36]. The ability to accurately detect and classify leaves is essential for understanding plant growth patterns, identifying diseases, and optimizing crop management practices. Researchers can track the growth and development of plants over time by analyzing the size, shape, and color of leaves, which provides insights into the health and vigor of plants. Leaf classification helps in identifying diseases that affect plants by analyzing distinct symptoms on leaves, such as discoloration or spots. By accurately classifying these symptoms using computer vision techniques, researchers can quickly diagnose plant diseases and take appropriate measures to prevent their spread. Hence the proposed Model as shown in Fig. 3, for the recognition of leaves in a video is the same for both the Training and Testing Phases but in the testing instead of storing to the Knowledge base it compares with the knowledge base for recognition. Furthermore, leaf detection and classification are instrumental in environmental monitoring efforts, as leaves act as natural indicators of air pollution levels since they directly interact with the atmosphere. By analyzing the condition of leaves in different regions or near industrial areas using video analysis techniques, scientists can assess air quality levels more efficiently. The detection and classification of leaves in a video is an important research area in the field of computer vision, and video analysis is a powerful tool here as it can provide temporal information about leaves. There are a lot of research opportunities in this field, some of which are mentioned below:

Data Acquisition

Most researchers focus only on image samples, which are collected manually. It is a tedious and time-consuming job, but it is very much needed in deep learning methods which require considerably large data samples for process. Large amounts of data can be acquired through video and generate required image samples in a few minutes by processing them using different image processing algorithms automatically.

Effective Sample and Feature collections

by using image processing techniques on a video frame to re-construct the samples and features as full or partially due to cluttered backgrounds, leaf occlusion frames/ images

3-D Feature building and extraction

the researchers are considering one side of the leaf (the front side) and ignoring other side features, 3d characteristics like the front, back, and thickness of a leaf may be useful in decision-making. So 3d features can be built by tracking video frame transmissions automatically or with minimal human interaction,

N-D Features / Metadata modeled

Along with 3d features, add present visual environment details like Forests, Deserts, Mountain regions, water surface regions, other associated plant parts, and some temporal information. which may play a crucial role in building an effective computational recognition system and also more generalized recognition system. This reduces the classification domain category selected based on the details(metadata). This can be achieved by video analysis, video frame tracking, and appropriate AI Techniques.





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Challenges and Future Directions

Challenges: Leaf detection and classification in videos are challenging tasks for the following reasons even with some of the challenges in images:

Video motion: The motion of leaves in videos can make it difficult to track and classify them accurately when they cross the speed limit of the camera. Despite the recent advances in deep learning-based leaf detection and classification, there are still some challenges that need to be addressed [35, 40]. One challenge is that deep learning models can be computationally expensive to train and deploy. Another challenge is that deep learning models can be susceptible to over fitting, especially when trained on small datasets. Future research in leaf detection is likely to focus on the following areas: Developing more robust and accurate deep-learning models for leaf detection and classification. Addressing the challenges of background clutter, varying illumination conditions, and occlusions. Developing real-time leaf detection methods with the help of 3d Features for recognition systems

CONCLUSION

In conclusion, deep learning has emerged as a powerful tool for leaf detection and classification [29, 31]. However, challenges remain in aspects like model efficiency, robustness, and real-time usage [35, 40]. Continued research should concentrate on translating accuracy in controlled settings to real-world applications [34, 37]. Overall, there are still gaps between experimental demonstrations and reliable field deployment that necessitate rigorous validation [36, 14].

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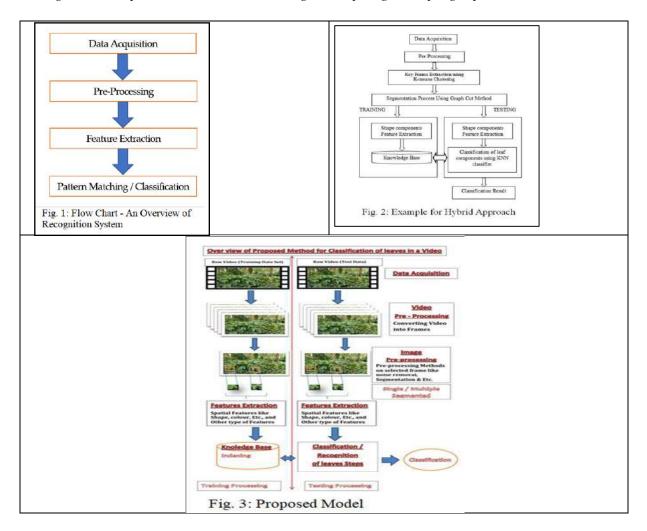
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RESEARCH ARTICLE

A Data Hiding Method Using Reduced Difference extension to develop the Stego Quality

Sumanth S1* and Siddarama S2

¹Associate Professor, Department of Computer Science and Applications, Government College for Women, Kolar – 563 101, Karnataka, India.

²Head, Department of Computer Science and Applications, Sri Bhagawan Mahaveer Jain College, Robertsonpet, K.G.F. - 563 122, Karnataka, India.

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*Address for Correspondence

Sumanth S

Associate Professor,

Department of Computer Science and Applications,

Government College for Women,

Kolar - 563 101, Karnataka, India.

Email: sumanth81s@gmail.com/dr.sumanth.s.81@gmail.com



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ABSTRACT

Because of the issues connected with network usage restrictions violations and unauthorized access to public networks, protecting secret data has taken on increased importance within network security. In response to these challenges, various data protection methodologies, including steganography, have been created. Steganography is concerned with securely conveying secret messages over public networks by hiding them within digital files. This study presents a unique steganographic approach based on reduced difference expansion, with the goal of reducing the magnitude of differences before embedding secret material. This study's experimental results show higher stego image quality, as indicated by a Peak Signal-to-Noise Ratio(PSNR) of 44.17 dB, a significant improvement over existing approach.

Keywords: Network infrastructure, national security, data protection, steganography, difference expansion

INTRODUCTION

The omnipresent demand for file sharing among users endures in the modern era of information technology, overcoming temporal and spatial boundaries. Despite its extensive use, the traditional use of public networks for such communications is fundamentally insecure. Because of the inherent vulnerability of this infrastructure, data traveling across the internet is vulnerable to interception and illegal alteration. The integrity and secrecy of the communication are jeopardized if an altered message reaches the intended recipient. As a result, data protection





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during the transmission phase becomes critical. One approach to achieving this safeguarding is by implementing data hiding techniques [1 - 3], wherein confidential information is concealed within diverse multimedia formats, including images, videos, audio, or text. Data hiding encompasses several approaches, such as watermarking, cryptography, and steganography. Different forms of covers have been utilized in steganography to maximize the confidentiality of the hidden data [4 - 6]. In the area of digital media steganography, establishing excellent visual quality of the stego medium while keeping a significant embedding capacity poses a significant difficulty [7]. The pronounced distortion introduced to the original media raises suspicions of attacks such as steganalysis [8 - 10] regarding the steganographic nature, potentially prompting adversaries to exert concerted efforts to compromise the embedding algorithm. Owing to the visual similarity between original and stego media, unauthorized interceptors may remain oblivious to the covert transmission of a concealed message. Typically, following the embedding of a secret message into the cover, the resultant stego medium traverses a public network to its destination, where the concealed message and cover are subsequently extracted and reconstructed. Researchers have proposed many data hiding approaches in the existing literature to keep the better quality of stego media while accommodating a defined embedding capacity, as proven by publications such as [11] and [12]. Furthermore, in [13], a spatial domain technique for steganography of digital images based on Difference Expansion (DE) was presented, utilizing redundancy information between pixels for secret message concealment.

Another approach, outlined in [14], involves developing a methodology affording high embedding capacity with minimal distortion in host images, employing RGB images, and inserting secret bits based on pixel block classification. Nevertheless, steganography in digital images presents a high trade-off gap between the stego image quality and the payload size. Several research works [15 - 17] have recommended using various combinations of mathematical functions to improve the positions and values of the pixels in the stego picture to address the issue of stego image distortion caused by concealment of the high payload size in the spatial domain. Recent work, however, indicates the necessity to continue working on decreasing the trade-off between stego quality and payload size. In this paper, we present a unique steganographic architecture that combines difference expansion with reduced difference expansion, with the goal of intentionally reducing the magnitude of differences before embedding hidden bitstreams into the cover's pixels. The primary goal of this research is to improve the quality of the generated stego image following data embedding. This manuscript is structured into five sections. Section I provides a comprehensive overview of the research problems under consideration. Section II presents an in-depth exploration of the existing literature on data hiding, mainly focusing on the difference expansion method. Section III elucidates the proposed method, detailing its conceptual framework. Moving on to Section IV, the evaluation of the method and its corresponding results are expounded upon, offering a comparative analysis with preceding methodologies. The conclusion of this work is included in Section V.

Related Work

Given the various technology developments integrated into digital image steganography in recent years, significant performance gains have been made. As a notable result, intelligent algorithms capable of securing confidential information have emerged. These algorithms are classified into two types based on their application domain: spatial domain techniques and transform (frequency) domain techniques. In the spatial domain, the concealment of confidential data involves direct manipulation of values of the pixels within the carrier to attain the wanted enhancement [13], ofparamount significance in contemporary digital image steganography are techniques such as difference expansion [13], expanded difference expansion [11], and pixel value modification (PVD) [12], all prominently situated within the spatial domain. These methodologies find prevalent usage when direct modification of fixed values of the pixels in an image is deemed necessary. Notably, they excel in achieving a concealment size; however, they occasionally exhibit susceptibility to compromising the stego image's quality. Among these techniques, difference expansion has emerged as one of the most renowned spatial domain algorithms. Its efficacy lies in concealing secret data by expanding the calculated values of the differences among pixels, contributing significantly to spatial domain steganographic methodologies. In work by researchers [13], an innovative reversible data hiding method was introduced, focusing on concealing data within images. This method was designed to





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concurrently uphold the quality of the resulting stego image and the payload capacity. Employing the binary-block embedding technique facilitated the concealment of data within the neighboring pixels' differences. Notably, a novel security key design approach was incorporated to fortify the algorithm against data loss, brute-force attacks, noise interference, and differential attacks. A noteworthy characteristic of this scheme lies in its complete reversibility, allowing for the restoration of embedded secret data independently and without any loss. This attribute enhances the utility and reliability of the proposed algorithm in the context of reversible data hiding within digital images. The work presented in [18] introduced a distinctive approach involving concealing secret data within two layers to augment load capacity. This is achieved through a security system integration that combines AES cryptography with image steganography, employing cross-division and additive homomorphism. The objective is to establish a novel reversible data hiding scheme tailored for encrypted images, thereby fortifying privacy protection in multimedia applications. The adoption of a homomorphic technique in this endeavor is particularly noteworthy. This choice is rooted in the capability of homomorphic operations to facilitate computations with encrypted data, thereby avoiding their expansion. As a result, the suggested scheme's overall embedding capability improves.

The embedding technique entails changing the cover image with a cryptography key and hiding the bits within the altered original image with the secret bits concealment key. To ensure reversibility, the strategy includes histogram shifting. Notably, data extraction occurs independently of image decryption at the receiver's end. This means that extracting hidden bits can be done before or after decrypting the altered image, giving the data extraction technique flexibility and variety. In a study outlined in [12], the authors introduced a method for hiding data by examining the differences between neighboring pixels after arranging them in a particular order. They leveraged the neighborhood of pixels, following a paradigm called pixel value ordering (PVO). Their approach involved sorting all the pixels in an image in ascending order and then pairing them off. Subsequently, they conducted a subtraction operation between these neighboring pixel pairs, using the results to conceal secret data. Notably, their technique does not use all the computed differences; instead, it customizes which pixels can hide data based on the values of these differences. They specifically choose pixels with differences of less than one for data embedding, resulting in a suboptimal payload capacity. As a result, further research is warranted to enhance their method's payload capacity. Using the knowledge gained from previous steganographic methods, this paper aims to provide a new steganographic technique that improves the quality of the stego image even when accommodating a large payload size. As cover images, general-purpose photographs from [19] are used.

PROPOSED METHODOLOGY

This section describes the steps taken for data embedding and extraction. We illustrate in Fig. 1 a flowchart for data embedding and in Fig. 2 a flowchart for data extraction. To shed much light on our method, we give a step-by-step description of the embedding process in Section III. A, and we present the pseudocode in Algorithm I for data extraction.

Data Embedding Steps

Step 1: Load the cover image (Cover).

Step 2: Arrange the pixels of the cover image into pairs-based blocks (1×2) .

Step 3: Compute the difference (h) between adjacent pixels (e0, e1) within each block using (1).

$$h = e1 - e0$$
 (1)

Step 4: Determine the block's characteristics based on the value of h. A block is considered "Changeable" if the difference is within the range of [-2, 7] ($h \le -2$ and $h \le 7$), "Expandable Non Reduced Difference Expansion (RDE)" if the block's difference is equal to 0 or 1 (h = 0 or h = 1)

Step 5: For the class of "Expandable RDE" blocks, we consider the difference values greater or equal to 2 and the ones less or equal to 7 ($h \le 2$ and $h \le 7$).





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Step 6: Create a location map with the same dimension as the cover image to store the block's characteristics. For "Unchangeable" blocks, assign the value of -1. For "Expandable Non-RDE" blocks, assign the value of 0. For "Expandable RDE" blocks, assign the value of 1.

Step 7: Conceal the secret data (s) in the calculated differences and obtain the new differences (*h*') using (2). We embed secret data directly for "Changeable" and "Expandable Non-RDE" blocks.

$$h' = 2 \times h + s \tag{2}$$

For "Changeable" and "Expandable" blocks, use a logarithmic function of base two to reduce the difference before we expand and embed the secret data using (3) to obtain a new reduced difference (h'').

$$h'' = h - (2[Sog2(h)] - 1 + log2(h\sqrt{h}) h)$$
 (3)

Step 8: Use the differences obtained in steps 6 and 7 to embed the secret data in the pixels of the cover image (ei) to obtain (ei"), the new pixel for the stego image using (4) and (5).

$$ei' = ei + h' \tag{4}$$

$$ei' = ei + h'' \tag{5}$$

Step 9: Construct the stego image and return the location map used in the secret data extraction process.

Data Extraction Process

This sub-section gives the details in pseudocode of how to extract the data in Algorithm 1. It is important to note that we can successfully extract the cover image and the secret data with the proposed method.

Algorithm 1: The pseudocode for the extraction process

Procedure extraction_process takes in parameters: stego_image, location_map, block_size

Initialize cover_image as a copy of stego_image

Initialize payload as an empty list

For each i in the range from 0 to the height of the location_map, stepping by block_size[0]

For each j in the range from 0 to the width of the location_map, stepping by block_size[1]

If the block does not exceed the image boundaries

Extract the block from the stego image and convert it to int16

Determine the block_type from the unique value in the corresponding location in location_map

If block_type is 0, continue to the next iteration

Else

If block_type is 1

Compute the difference between the first two pixels in the block

Append the least significant bit of the difference to payload

Compute the original difference by floor division of the difference by 2

Update the second pixel in the block in cover_image with the sum of the first pixel in the block and the computed difference

Else

Compute the difference between the first two pixels in the block

Append the least significant bit of the absolute difference to payload

Compute the original difference using Equation 2.

Update the second pixel in the block in cover_image with the sum of the first pixel in the block and the computed difference

Return cover_image, payload

End Procedure

RESULTS AND DISCUSSIONS

In this section, we report the PSNR results in Fig. 3. Moreover, we present a comparative view of our results to those obtained in [Maurice] in Table 1. Table 1 thoroughly examines PSNR values, comparing the method delineated in





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[11] with our novel proposed approach. The assessment encompasses a range of cover images, namely Abdominal, Leg, Head, Hand, and Chest, and investigates three distinct payload sizes: 5 kilobits, 10 kilobits, and 20 kilobits. The PSNR values, measured in decibels (dB), serve as crucial metrics to evaluate the quality of stego images produced by both methods under varying payload conditions. This table1 allows for a detailed analytical view of the performance variations between the existing method and our approach across different cover images and payload sizes, highlighting the consistent superiority of our proposed method in maintaining high-quality stego images. Figure 3 illustrates an analytical portrait of our method for payload sizes (in kb) and the corresponding PSNR values for distinct cover images. Cover images include Abdominal, Leg, Head, Hand, and Chest, with payload sizes ranging from 5 kb to 20 kb. The PSNR values highlight the quality of stego images produced by our proposed method at different payload levels and across diverse cover images. This figure emphasizes how the proposed method maintains image quality while accommodating varying payload sizes, demonstrating its versatility and efficacy in concealing data within digital media.

CONCLUSION

In today's technology landscape, digital photographs have become prevalent elements used for a variety of applications. Digital photographs, because of their ubiquitous use, serve an important role in facilitating the transmission of secret information across public networks. In our study, we present a unique way for increasing the confidentiality of secret data. This method makes use of developments in difference expansion techniques to disguise secret information within digital photographs. This study addresses the significant compromise in stego image quality inherent in spatial domain image steganography. While previous research efforts addressed this issue, a clear trade-off between PSNR and payload size persisted. In comparison to existing approaches, our solution uses an expanded difference expansion to improve performance. Our method's results show a promising advantage over previously described ways, allowing secure transmission of large-sized payloads within digital images while keeping the quality of the cover image.

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Table 1. Comparison of our results with the existing method in maurice [11]

	PSNR (dB)							
Cover Image]	Proposed Method						
intage	5kb	10kb	20kb	5kb	10kb	20kb		
Abdominal	40.87	38.18	38.17	41.58	38.67	38.66		
Leg	40.62	38.93	38.83	41.14	39.8	39.65		
Head	34.74	32.72	32.73	35.83	33.54	33.55		
Hand	40.65	38.03	38.03	41.44	38.5	38.5		
Chest	42.46	39.15	39.05	44.17	40.23	40.04		





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Average 39.868 37.402 37.362 40.83 38.15 38.08

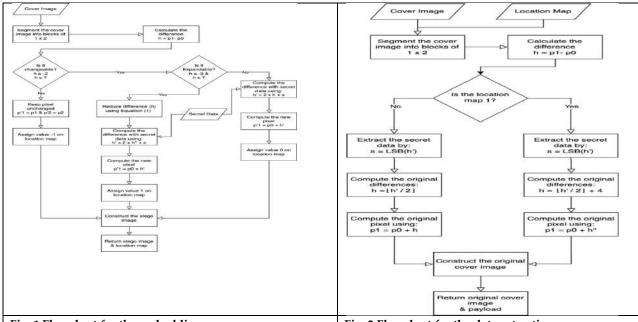


Fig. 1 Flowchart for the embedding process.

Fig. 2 Flowchart for the data extraction

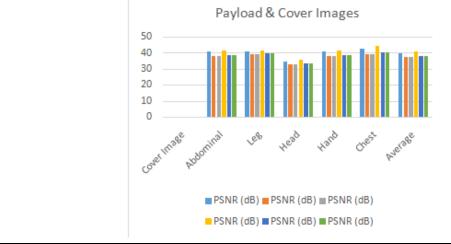


Fig. 3 The Obtained PSNR with the Proposed Method





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RESEARCH ARTICLE

Recognizing Handwritten Digits Using Deep Learning Algorithms

N. Sumathi^{1*}, Pranesh. K.N² and Siddarth A.K²

¹Associate Professor, Department of Information Technology, Sri Ramakrishna College of Arts & Science Coimbatore, Tamil Nadu, India.

²PG Student, Department of Information Technology, Sri Ramakrishna College of Arts & Science Coimbatore, Tamil Nadu, India.

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*Address for Correspondence

N. Sumathi

Associate Professor, Department of Information Technology, Sri Ramakrishna College of Arts & Science Coimbatore, Tamil Nadu, India. Email: sumathi.n@srcas.ac.in



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ABSTRACT

Humans now rely on technology more than ever before, and deep learning and machine learning algorithms enable us to do everything from classify objects in photos to add music to silent movies. Similarly, one of the important fields of study and advancement with a seemingly endless supply of potential applications is handwritten text recognition. Handwriting recognition (HWR), alternatively referred to as handwriting Text Recognition (HTR), refers to a computer's capacity to read and comprehend legible handwriting input from many sources, including paper documents, images, touch displays, and other devices. In this paper, it appears that we used Support Vector Machines (SVM), Multi-Layer Perceptron (MLP), and Convolution Neural Network (CNN) models to accomplish handwritten digit recognition with the aid of MNIST datasets. Our major goal is to determine which model is optimal for digit recognition by comparing the execution times and accuracy of the models mentioned above.

Keywords: Multi-Layered Perceptron (MLP), Convolution Neural Network (CNN), MNIST datasets, Deep Learning, Machine Learning, Handwritten Digit Recognition, Support Vector Machines (SVM).

INTRODUCTION

A computer's capacity to identify handwritten letters and numbers from a variety of media, including pictures, documents, touch screens, and more, and group them into ten preset categories is known as handwritten digit recognition (0-9). In the realm of deep learning, this has been an endless source of research. Numerous tasks, such as





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sorting mail, processing bank checks, and recognizing license plates, need the use of digitization [1-7]. We have several difficulties in handwritten digit recognition since various people write in different ways, and this type of writing is not optical character recognition. To achieve the goal of handwritten digit recognition, this study offers a thorough comparison of several machine learning and deep learning techniques. Such as Convolutional neural networks, multilayer perceptrons, and support vector machines. Based on the algorithms' accuracy, mistakes, and testing-training times, a comparison is made between them, supported by charts and graphs created using matplotlib for visualization. The accuracy of any model is important because more accurate models yield better outcomes. Real-world applications are unsuitable for models with low precision. Ex: High accuracy is crucial for an automated bank check processing system that can identify the amount and date on the check. It is not ideal if the system detects a digit wrongly since it might cause significant harm. For this reason, a highly accurate algorithm is needed in many practical applications. Therefore, we are presenting an accuracy comparison of several methods so that the most accurate algorithm with the lowest probability of mistakes may be used in a variety of handwritten digit recognition applications[8]. For handwritten digit recognition, this article offers a fair overview of machine learning and deep learning techniques including SVM, CNN, and MLP. Additionally, it provides with the knowledge of which algorithm works best for digitrecognition[9-12]. For a more equitable knowledge of the three algorithms, we will first address the technique and implementation of the relevant work that has been done in this subject in later portions of this study. Subsequently, itshowcases the outcome and conclusion, supported by the research conducted for this paper. Additionally, it will provide with some possible future developments in this subject. The references and citations utilized are included in the latter portion of this work.

RELATED WORK

Tan and Le proposed EfficientNet, which provides valuable insights into optimizing CNN architectures for improved performance [13]. He et al. introduced ResNet, a groundbreaking CNN architecture that enables the training of extremely deep networks, leading to superior performance in image recognition tasks. ResNet has become a cornerstone in CNN research and has been widely adopted in various applications [14]. Sun et al. revisited the importance of data in the deep learning era and emphasized the need for large-scale, diverse datasets to train robust deep learning models effectively. [15]. Liu et al. proposed DARTS+, an improved version of Differentiable Architecture Search (DARTS), which addresses the instability and inefficiency issues associated with previous methods. By introducing early stopping and weight decay techniques, DARTS+ achieves more stable and reliable architecture search results [16]. Zhang et al. introduced Squeeze-and-Excitation (SE) networks, which allows CNNs to focus on informative features while suppressing irrelevant ones, leading to improved performance in image classification tasks [17]. Pham et al. proposed an efficient method for Neural Architecture Search (NAS) by introducing parameter sharing among child models. This approach significantly reduces the computational cost of architecture search, making it feasible to explore a large search space efficiently [18].

Cubuk et al. introduced Auto Augment, a data augmentation technique that automatically learns augmentation policies from the training data. By dynamically applying a combination of augmentation transformations, Auto Augment improves model generalization and performance. This work highlights the importance of data augmentation in training robust CNN models [19]. Vaswani et al. proposed the Transformer model, which utilizes self-attention mechanisms to capture long-range dependencies in sequential data [20]. Simonyan and Zisserman introduced VGGNet, a CNN architecture known for its simplicity and effectiveness. VGGNet consists of multiple convolutional layers with small filter sizes, stacked on top of each other[21]. The existing handwritten digit classification system utilizes a Support Vector Machine (SVM) algorithm to compare pixel values of input images with known digit representations, ultimately identifying the closest match and outputting the corresponding digit. Despite its simplicity, the system faces challenges in accuracy and scalability. Accuracy is compromised by variations in input image quality, while scalability is hindered by increasing dataset size, leading to computational complexity. Additionally, the pixel-level comparison may fail to capture intricate features in handwritten digits, resulting in suboptimal classification accuracy, especially for complex inputs. Unlike the existing system based on Support Vector Machine (SVM), CNNs offer superior accuracy and robustness to variations in input image quality.





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PROPOSED METHODOLOGY

The proposed system for handwritten digit classification adopts a Convolutional Neural Network (CNN), a specialized deep learning algorithm tailored for image classification tasks. Unlike the existing system based on Support Vector Machine (SVM), CNNs offer superior accuracy and robustness to variations in input image quality. By leveraging CNNs, the proposed system can effectively capture intricate features in handwritten digits, leading to enhanced classification accuracy. Additionally, CNNs can be trained on larger datasets containing known digit representations, further improving their accuracy and generalization capability. Overall, the proposed CNN-based system offers a more advanced and effective approach to handwritten digit classification, promising superior performance and reliability compared to the existing SVM-based system.

Model Architecture

Design the Model Architecture: The architecture of the classification model is designed, considering options such as a Multi-Layer Perceptron (MLP) with one hidden layer or a Convolutional Neural Network (CNN). The chosen architecture should be capable of effectively learning and classifying handwritten digits from the MNIST dataset. The Architectural layer of CNN shown in figure 1 depicts the working process of the recognizing system.

Techniques

- 1. MLP: A simple feed forward neural network with one hidden layer. The number of 10 neurons in the hidden layer, activation functions, and output layer size are defined based on the requirements of the classification task.
- 2. CNN: A more complex architecture comprising convolutional and pooling layers followed by fully connected layers. CNNs are well-suited for image classification tasks and can capture spatial hierarchies of features in the input images.

Model Training

Train the Model

The model is trained by feeding training images and corresponding labels into the neural network. During training, the model adjusts its parameters through backpropagation, where gradients of the loss function with respect to the model parameters are calculated and used to update the weights of the network. This iterative process enables the model to learn to correctly classify digits based on the input images.

Techniques

- 1. Back propagation: Back propagation is a key technique used in training neural networks. It involves propagating the error backwards from the output layer to the input layer, adjusting the weights of the network to minimize the error. This process is repeated iteratively until the model converges to a satisfactory solution.
- 2. **Optimization Algorithms (Adam):** Optimization algorithms like Adam are employed to efficiently minimize the loss function during training. Adam combines the advantages of two other popular optimization techniques, AdaGrad and RMSProp, by maintaining separate learning rates for each parameter and adapting them based on past gradients and squared gradients.

Data Loading and Preprocessing

- 1. Load MNIST Dataset: Use TensorFlow or Keras to load the MNIST dataset, containing handwritten digit images and corresponding labels.
- 2. Normalize Pixel Values: Apply min-max normalization to normalize pixel values of the images to the range [0, 1]. This normalization ensures uniformity in input feature scales, aiding in efficient training and model convergence.
- 3. Convert to Grayscale: Convert the images to grayscale to remove color information, simplifying the input data while retaining relevant features necessary for digit classification.
- 4. Resize Images: Resize the images to a desired dimension, typically 28x28 pixels. Resizing ensures uniformity in input size across all images.





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IMPLEMENTATION

We have utilized three distinct classifiers to compare the algorithms based on working accuracy, execution time, complexity, and the number of epochs (in deep learning methods):

- 1. Convolutional Neural Network Classifier;
- 2. ANN Multilayer Perceptron Classifier;
- 3. Support Vector Machine Classifier.

The implementation of each method has been covered in depth below, along with the analysis's flow to produce an accurate and seamless comparison.

PRE-PROCESSING

The first stage of machine and deep learning is called pre-processing, and it aims to enhance the input data by removing unnecessary redundancies and contaminants. All of the photos in the dataset were molded into 2-dimensional images, or (28,28,1), in order to simplify and deconstruct the input data. Since the picture pixel values varied from 0 to 255, we normalized the pixel values by dividing the dataset by 255.0 and converted the dataset to 'float32.' This allowed the input features to range from 0.0 to 1.0. The y values were thentranslated into zeros and ones using one-hot encoding, which made each number categorical. For instance, an output value of 4 would be transformed into an array of zeros and ones, or [0,0,0,0,1,0,0,0,0,0]

SUPPORT VECTOR MACHINE

Scikit-learn's SVM accepts as input sample vectors that are sparse (any scipy.sparse) or dense (numpy.ndarray and converted to that by numpy.asarray). SVC, NuSVC, and LinearSVC in scikit-learn are classes that can sort data into several classes. Using a linear kernel that was constructed with the aid of LIBLINEAR, we employed LinearSVC in this research to classify MNIST datasets [22]. The implementation has made use of several scikit-learn packages, including NumPy, matplotlib, pandas, Sklearn, and seaborn. Prior to loading and reading the CSV files using pandas, we will first obtain the MNIST datasets. Following that, several samples were plotted, and the data was converted into a matrix, normalized, and scaled. In the end, we developed a linear SVM model and a confusion matrix to assess the model's accuracy.

MULTILAYERED PERCEPTRON

The application of feed forward artificial neural networks, or multilayer perceptrons, for the recognition of handwrittendigits is carried out by utilizing the Keras module to build an MLP model of the sequential class and add corresponding hidden layers with distinct activation functions to receive an image with a pixel size of 28 x 28 as input. We added a dense layer with various requirements and drop out layers after building a sequential model. We employed an output layer with 10 units (i.e., the total number of labels) in a neural network with 4 hidden layers. There are always 512 units in the hidden layers. The 784-dimensional array created from the 28×28 picture serves as the network's input. We constructed the network using the Sequential model. By adding the necessary layer one at a time, we may simply stack layers in the sequential model. Since we are creating a feed forward network in which all of the neurons from one layer are connected to the neurons in the layer before it, we utilized the Dense layer, also known as a completely connected layer. In addition to the Dense layer, we included the ReLU activation function, which is necessary to give the model non-linearity. By doing this, the network will be able to learn non-linear decision limits. Since the problem involves multiclass classification, the final layer is a softmax layer.

CONVOLUTIONAL NEURAL NETWORK

Keras is used in the development of Convolutional Neural Network for handwritten digit recognition. It's an open-source neural network library used for deep learning model construction and implementation. The Convolutional Neural Network (CNN) architecture, depicted in figure 2, is a powerful deep learning model widely employed for image recognition tasks. The network comprises several key components arranged in a sequential manner. At the





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beginning lies the Input Layer, where raw image data is ingested. Subsequently, Convolutional Layers process the input through learnable filters, detecting features like edges and textures. Following each convolution operation, Activation Functions, typically ReLU, introduce non-linearity to the network, aiding in capturing complex patterns. The feature maps produced are then subjected to Pooling Layers, which down sample the spatial dimensions while retaining crucial information. The Fully Connected Layers aggregate these high-level features, leading to the final Output Layer, where predictions or classifications are made. Throughout training, a Loss Function measures the disparity between predicted and actual outputs, guiding optimization via back propagation. This process is facilitated by Optimization Algorithms such as Stochastic Gradient Descent, ensuring continual refinement of the network's parameters. This CNN framework enables robust and efficient learning, making it a cornerstone in modern image analysis and classification tasks.

EXPERIMENTAL RESULTS

Handwritten character recognition is a broad field of study with well-defined implementation strategies already in place. These strategies include popular algorithms, large learning datasets, feature scaling, and feature extraction techniques. The NIST dataset is made up of two NIST databases: Special Database 1 and Special Database 3. A portion of the NIST collection is the MNIST dataset, or Modified National Institute of Standards and Technology database. Special Database 1 and Special Database 3 comprise entries penned by high school students and US Census Bureau staff, respectively. The MNIST dataset comprises 70,000 handwritten digit pictures in a 28x28 pixel bounding box with anti-aliasing (60,000 for the training set and 10,000 for the test set). Each of these pictures has a matching Y value that indicates the digit. We examined the accuracy and execution time of the three algorithms-SVM, MLP, and CNN-after putting them all into practice using experimental graphs to ensure clear comprehension. All the previously mentioned models' training and testing accuracy have been considered. After running every model, we discovered that SVM achieves the best accuracy on training data, whereas CNN achieves the best accuracy on testing data. To learn more about how the algorithms function, we have also compared the execution times. An algorithm's running time is typically influenced by the quantity of operations it has completed. To obtain the desired result, we have trained our SVM models using norms and our deep learning model for up to 30 epochs. CNN accounts for the most running time, whereas SVM requires the least amount of time to execute. Each model's accuracy is shown in this Table 1. Additionally, we showed how deep learning models improved accuracy and decreased error rate in relation to the number of epochs by visualizing their performance measure. Drawing the graph is important because it helps us determine whether to apply an early stop and prevent over fitting, which occurs when accuracy changes steadily over a period of epochs. precision measures the model's ability to make accurate positive predictions while minimizing false positives as shown in the figure 3.

CONCLUSION AND FUTURE ENHANCEMENT

Using MNIST datasets, this research study constructed three models for handwritten digit recognition: Support Vector Machine, Multi-Layer Perceptron, and Convolutional Neural Network. It evaluated them based on the number of epochs (in deep learning techniques), working accuracy, complexity, and execution time to determine which model was the most accurate and came to the following conclusion: Because SVM is a fundamental classifier, it performs faster than most other algorithms and, in this instance, yieldsthe greatest training accuracy rate. However, because of its simplicity, it is unable to identify complicated and ambiguous pictures with the same level of accuracy as MLP and CNN algorithms. When time is considered, it is discovered that the MLP model produces nearly the same accuracy rate as the CNN model in a very efficient amount of time. However, the only disadvantage is that it is unable to sustain the classification rate in complex images for extended periods of time because it is unable to comprehend the spatial relationships between image pixels to the same extent as CNN. The highest accurate results for handwritten digit recognition were obtained using CNN; the only disadvantage is that it required exponential





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processing time. Due to a particular model's limitations, extending the number of epochs without altering the algorithm's configuration is pointless. It has been observed that the model begins to overfit the dataset beyond a predetermined number of epochs, which results in biased predictions. Thus, it confirms that, when compared to SVM and MLP, CNN has the highest accuracy rate for any kind of prediction issue including image data as an input. The possibilities for developing applications based on deep and machine learning techniques are essentially endless. In the future, it will be possible to solve many issues by working on denser or hybrid algorithms that use more manifold data than the current collection of methods.

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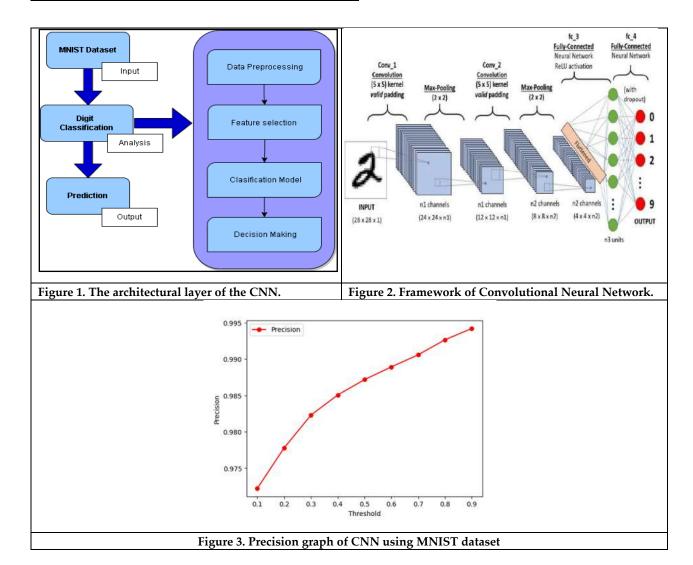


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Table 1: Comparison Analysis of Different Models.

S.NO	MODEL NAME	ACCURACY		
1	SVM	94.65%		
2	MLP	98.85%		
3	CNN	99.56%		







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REVIEW ARTICLE

Security, Trust, and Privacy of Information Sharing **Networking Sites: A Systematic Literature Review**

V. Padmaja¹, MuniKrishnappa AnilKumar², Sirigiri Srikanth^{3*}, Pocha Sri Chandana³.

¹Professor, M.S. Ramaiah Institute of Management, Bengaluru, Karnataka, India.

²Assistant Professor, M.S. Ramaiah Institute of Management, Bengaluru, Karnataka, India.

³PGDM 1st year, M.S. Ramaiah Institute of Management, Bengaluru, Karnataka, India.

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*Address for Correspondence Sirigiri Srikanth

PGDM 1st year,

M.S. Ramaiah Institute of Management,

Bengaluru, Karnataka, India.

Email: sirigirisrikanth231139@msrim.org



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ABSTRACT

The influx of netizens using social media platforms, sharing of information online has become the norm, and there is a constant threat to security and data privacy concerns. Though much research has been done on privacy issues emerging in social media interactions, very limited systematic literature review has been done to get a comprehensive understanding of the issue. This study has been undertaken through a systematic literature review which highlights the challenges and opportunities related to security, trust, and privacy in social networking sites. The study includes three social networking sites namely Facebook, Twitter and Instagram, and has identified potential strategies, technological solutions, and policy implications to mitigate risks and foster a safer and more trustworthy social media environment, which can be beneficial to users and platform developers. The results of the study have significant implication to academicians, policy makers, and industry on current understanding of the problem area. The study will enhance a comprehensive understanding of the complex dynamics like security threats, privacy concerns, and trust issues inherent in the context of social media information sharing, with a focus on security, trust, and privacy. It is a significant endeavor with potential real-world implications.

Keywords: Security, Trust, Privacy, Information, Social networking sites.

INTRODUCTION

With the advent of social networking sites, individuals were seen voluntarily disclosing personal information in various forms, which raises especially important questions for individual privacy and civil liberties (Benson et al.,





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2015). Social media networking sites serve as virtual spaces where individuals, groups, and organizations can connect, share information, and interact with each other. They facilitate communication, collaboration, and community-building across geographic boundaries (Gupta & Dhami, 2015). Social networking sites have changed from a social experiment to a way of interpersonal communication (Papaioannou et al., 2021). These platforms differ from one another and offer unique features to their users, with wide variety of options available on social networking sites. We have created and maintained virtual profiles, which are digitalized and contain different sets of information about their users for this purpose. Nowadays, the cause of excessive information sharing of information online in the digital age has its advantages and disadvantages, the whole depends on how you present yourself to others in online. Digitalization of data raises issues such as security, trust, privacy of sharing personal information on online. These social network sites have a large user database that provides users with information across the Globe. Thus, the information that we share, and shared on these platforms may not always be safe and secure as we expect, and there are certain risks pertaining to the data disclosure. This paper helps to understand, analyze, and take action to reduce the sharing of our personal information on online. It has also educated the netizens about the awareness of information sharing on social media networking sites with some real-world implications. Social network administrators have taken several steps to ensure the security of users' information and privacy. There are several privacy and trust concerns that social network users need to know about (Papaioannou et al., 2021). The main concern of sharing the information is that the information is more visible and accessible where the issues pertaining to security, trust, and privacy arise. To ensure the privacy of users, social network administrators have implemented several security measures. For the purpose of this research, the following questions have been framed:

RQ1: is there any safety about the information that is being shared on social networking sites?

RQ2:what kind of awareness can be created amongst the netizens regarding information being shared online? The key challenges and issues discussed in this paper will ease the process of understanding and mitigating risks and thus systematically developing various dimensions of security, trust, and privacy.

METHODOLOGY

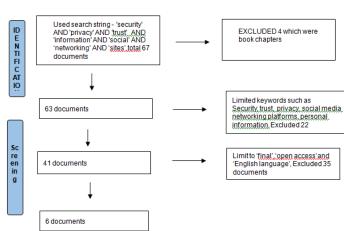
The study focuses on secondary data available in Scopus data base, inthe form of existing literature and research papers published on this topic. The Scopus database was filtered using the search string [TITLE/ABSTRACT] OR 'security' AND 'privacy' AND 'trust' AND 'social' AND 'media' AND 'networking' AND 'sites' [TITLE/ABSTRACT/KEYWORD], the operator used namely 'AND.' Further, articles were filtered using PRISMA.PRISMA is an evidence-based on minimum set of items for reporting systematic reviews and meta-analyses. PRISMA stands for Preferred Reporting Items for Systematic Reviews and Meta-analyses. The latest articles on security, privacy, and trust concerns of information sharing were taken into consideration while drafting this research paper. This is a Boolean Search of using a combination of "AND, OR" it produces accurate results while avoiding the irrelevant. Various cyber security websites such as AAG IT Services and Gitnux have been referred for statistics that has been used for actual data. A total of 67 documents (as on December 14th, 2023) were available in Scopus data base on the theme 'security, privacy, and trust' in social networking sites and six research papers were available in the context. The documents were downloaded and stored in the reference management software, Zotero. These papers have been further filtered based on 'document type', 'publication stage', and 'language'. Final filtration with six documents have been considered for the study. The discussion and findings have been framed after reviewing these six research papers.





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EXCLUDED 4 which were book chapters



Findings

PRISMA

As the social media hacking statistics show(AAG IT Services), any individual can be a target for any reason. Multibillion-dollar corporations can be attacked as well as individuals with modest incomes. Regardless of the identity of an individual, what platforms are used by an individual or how much wealth one has, anyone can be targeted. According to report of AAG IT Services (Jan 2024), 68000 users have enquired about remedies on Facebook when it was hacked, and even Instagram has got 36000 queries a month on the remedies for hacked accounts. As per the report of Gitnux, 530 million Facebook accounts are compromised by exposing their passwords, phone numbers, and account names. It is the same month and year for LinkedIn as well, where 750 million users i.e., 93% of total users resulted in the exposure of the client's phone numbers, emails, usernames, and geolocation records. As per the report compiled (January 2024) by AAG IT Services, a staggering 85% of Instagram accounts have been compromised, with an additional 25% of Facebook accounts falling victim to hijacking. These statistics are quite alarming, and highlight the urgent need for enhanced cybersecurity measures to protect individual's personal information and privacy online. Moreover, major corporations face relentless attempts by black hat hackers, with an average of 30 annual hacking incidents on their corporate social media accounts alone. The high-profile nature of some victims, from business tycoons to politicians, serves as a stark reminder of the pervasive threat posed by cybercrime in today's digital age. Real-time data breach cases can help to understand the stated objectives in this research.

Facebook data breach

Despite Facebook (META) being the giant in social networking sites, its app's developer Aleksandar Kogan, was allowed to gather data on users. In the end, Kogan amassed data from 87 million users, which he then gave to the political consulting business Cambridge Analytica in 2016. Cambridge Analytica utilized the data to create voter profiles and their political views to target and influence public opinion. Whistleblower Christopher Wylie of Cambridge Analytica claims that the company utilized this information to assist Donald Trump's campaign in influencing votes in the 2016 presidential election. Many have referred to this episode as a data breach. European and American regulators have forced CEO Mark Zuckerberg to appear before Congress, and the U.S.Federal Trade Commission and other international agencies are conducting an official inquiry into Facebook's privacy policies. This incident has raised doubts about the information netizens give to these networking sites and has emphasized the importance of being aware of what they post and who they believe.





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Twitter data breach

A significant security breach on Twitter in July 2020 resulted in the compromising of well-known accounts and Bitcoin fraud. Twitter itself confirmed that the intruders had gained access to Twitter's administrative tools so that they could alter accounts themselves and post the tweets directly. This made the platform suspend tweets from verified accounts until the issue was resolved. The law enforcement authority's investigation was made and there was a negative impact on the stock value of Twitter. People were worried about how secure their accounts were and how open the platform was to these kinds of assaults. The incident prompted inquiries over the efficacy of Twitter's security protocols. Twitter must put in place more security measures and be open and honest about the issue to win back user trust

Instagram data leak

A large database operated by a third-party company holding the confidential data of millions of Instagram influencers, celebrities, and brand accounts was uncovered by a security researcher in 2019. High-profile Instagram user's email addresses and contact details were made public. 49 million Records were exposed in this data leak. The database was linked to a marketing firm in India. People were concerned about the possible exploitation of their personal information, especially celebrities and influencers. The issue brought to light worries over Instagram user data access by other parties. Users' confidence in the security of Instagram was impacted by critics of the platform's data protection policies.

DISCUSSIONS

In today's digital age, where information is readily shared and accessed through social media networking platforms, security, trust, and privacy concerns have become paramount. Recent incidents such as the Instagram data scraping in 2021, TikTok's ongoing data privacy concerns, and Google's data breach in 2018 have brought to light about the vulnerabilities that users face, when sharing personal information on online. These events have scored the importance of robust security measures and increased transparency from social media companies to protect user data. The mishandling of user information on these platforms not only erodes trust between users and theplatform but also raises questions about the commitment of these companies to safeguard user privacy. As increased people rely on social media for communication, entertainment, and even business purposes, stringent protocols must be put in place to prevent unauthorized access to personal data. Social networking sites consist of virtual profiles that maintain ties and can be shared with others (*Papaioannou et al.*, 2021). A digital identity is formed through information that the individual shares in public and personal data disclosure is part of this process. Individuals come into contact, with information that compares them socially with others, such as photos, which can negatively affect their personal lives and privacy (*Gupta & Dhami*, 2015). In this digital age, it has majorly raised concerns about privacy, permission, and responsible data usage due to the vast amount of personal information collected and shared online. The ethical dilemma lies in striking a balance between privacy and tailored services.

Unauthorized data usage or repurposing can have unanticipated negative effects on public and commercial sectors. Some academicians argue that in a networked world, information privacy is no longer under the control of individuals but rests with organizations holding the information (*Benson et al.*, 2015). Moreover, privacy has evolved significantly due to technological advancements, social changes, and cultural shifts. As online interactions and personal information sharing become common, users' behavior on social media platforms has changed. Privacy concerns refer to a user's concerns about the possible loss of privacy due to voluntary or surreptitious information disclosure (*Chang & Liu*, 2023). The security, trust, and privacy concerns surrounding information sharing on social media networking platforms are more prevalent than ever. With the ability to personalize user experiences and perpetuate biases, these platforms can inadvertently create filter bubbles that influence user perceptions and exacerbate societal divisions. Additionally, the potential for governments and law enforcement agencies to leverage social media data for surveillance purposes raises fundamental questions about civil liberties and the right to privacy. Users seek transparency on data usage and platform operators, and companies with ethical practices gain





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more user trust. Social media platforms are popular for both social interactions and business transactions, presenting new challenges around information and their privacy (Benson et al., 2015). Furthermore, the wealth of personal information contained within social media profiles leaves users vulnerable to identity theft and impersonation. The ease of communication and anonymity provided by these platforms also open the door to cyber bullying and harassment, posing risks to user safety and well-being. To further our understanding of information security and privacy, when sharing data on social networking sites, we have carried out an experiment. To do this, we have made a few virtual accounts and posted some bogus information in them. We learnt that some intruders had acquired the material information and were sending us spam emails and links through various accounts that helps to easily track mobile phones and the information stored in them as well. Another significant instance involves cyber-stalkers who create fake profiles, pursue strangers, and gather their personal information for the purpose of data exploitation and extortion. These crimes have become so severe that even cybercrime can't control them to the full extent. The research further elaborates about the measures that must be taken by netizens and social media platform developers to mitigate the risks. The following measures can be takento mitigate the challenges faced by netizens:

By platform developers

Protect data with intrusion detection, secure authentication, and robust encryption. Update your software and security patches often to keep up with emerging danger. Utilize cutting-edge threat detection technology to promptly identify and address security problems. Give consumers more control over their privacy by giving them access to more options including evaluating app permissions, restricting data sharing, and modifying visibility settings. Keep an eye out for unusual activities, security breaches, and questionable conduct on social networking. Establish a robust incident response strategy event promptly, limit harm, and lower the possibility of user impact. Provide protocols for communicating about crises a reporting data breaches. Work together to share threat intelligence, best practices and helpful law enforcementwith social media platforms, cyber research, law enforcement, and other industry partners.

By users

Set distinct passwords for every social media platform you use. Employ capital, lowercase, and special characters. To generate and save distinct passwords for your accounts, use only a reliable password manager. If at all feasible, use two factor-authentication. Make sure you know who sent the message before clicking. Steer clear of abbreviated URLs that conceal dangerous links. When you get demands for private information, especially through unsolicited emails or messages, use caution. To confirm the authenticity of the request, get in touch with the sender. To manage who may see your information, relationships, posts, and profile on social media, you should periodically review and modify your privacy settings. Restrict how much personal information you post on social media. That too reliable contacts. Recognize typical cyberthreats, assaults, and strategies employed by cybercriminals, such as phishing scams.

Scope for further research

The research can be extended to a mixed-method approach by using both qualitative and quantitative data. Also, more databases can be referred to understand the dimensions of trust, privacy, and security to get a more comprehensive understanding.

CONCLUSION

The study highlights the importance of adopting a strategic and holistic approach to data privacy, security, and trust. By being more vigilant about the information we share in online and actively managing our privacy settings, individuals can better protect themselves from potential risks. Social media companies must also prioritize user data protection by implementing robust security measures and ensuring transparency in how they handle our





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information. Only through collective efforts can we effectively address the challenges of data privacy and ensure that all users are adequately protected in the digital landscape.

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RESEARCH ARTICLE

Detecting Deep Fakes: Leveraging Convolutional Neural Networks for **Fabricated Media Classification**

Prem Sagar*

Assistant Professor, Department of Computer Science St. Joseph's University, Bangalore - 27, Karnataka, India.

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*Address for Correspondence

Prem Sagar

Assistant Professor, Department of Computer Science, St. Joseph's University, Bangalore - 27, Karnataka, India. Email: prem.sagar_@sju.edu.in



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ABSTRACT

Deepfakes, synthesized media in which a person's likeness is swapped into a source image, are becoming increasingly harmful. Face swapping into intimate imagery infringes on personal privacy. Detecting deep fakes and differentiating them from real videos is therefore an important challenge. This research proposes a deep learning (Machine Learning) approach for deep fake detection. A Convolutional neural network model with Python and OpenCV is trained on a dataset of real and synthesized images to classify images as real or fake. Key features used by the models include inconsistencies in facial expressions, lighting, and skin textures that expose manipulated imagery. The work demonstrates deep learning's capabilities in identifying fabricated media and lays the groundwork for reliable deepfake classification. This technology will help curb the spread of misinformation and authentication of visual evidence as deep fakes grow more advanced and accessible.

Keywords: Deepfake, Machine Learning, Neural networking, Privacy, Manipulated imagery, convolutional neural network.

INTRODUCTION

Deepfakes, a combination of "deep learning" and "fake", refer to media generated or manipulated by artificial intelligence to depict events that never occurred or to falsify identities. Advances in generative adversarial networks and autoencoders have enabled creation of increasingly realistic deepfakes, spurring alarm over potential misuse for non-consensual pornography, financial fraud through identity theft, and disinformation campaigns. Detecting deepfakes has therefore emerged as a major challenge at the intersection of machine learning, computer vision, and information forensics.





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Convolutional neural networks (CNNs) have become essential in many artificial intelligence tasks involving visual data. As research progresses on deepfake classification, CNN architectures have emerged as promising solutions due to their capabilities in feature extraction. CNNs incorporate insights from processing in the human visual cortex and contain specialized layers optimized for spatial hierarchical feature maps. This equips them well to accurately classify real versus synthetic image or video samples. Though originally focused on the computer vision challenge of image recognition, CNNs now demonstrate success across natural language processing, medical imaging, and other applications. For deepfake detection specifically, CNN models show strength in learning robust discriminative representations to identify manipulated regions. While early detection methods focused on computer graphics artifacts, advanced techniques use convolutional neural network (CNN) architectures to learn robust visual features. However, the tools used to create deepfakes are getting better at making them look real and tricking detection systems. This back-and-forth competition between creating and spotting deepfakes means we need to improve how we design CNN models to keep up. This study aims to benchmark convolutional network architectures for deepfake classification based on face and scene representations. This paper carefully selects new datasets that combine computer graphics and neural synthesis techniques to cover distributional gaps. Using these datasets, experiments evaluate the accuracy and operational characteristics of the receiver and identify optimal model architectures, loss functions, and training regimes. This method and analysis can help forensic and fact-finding experts to authenticate media sources.

Deepfake Creation

Deepfakes leverage generative adversarial networks (GANs) to alter or synthesize visual content by modeling underlying data distributions. GANs comprise opposing generator and discriminator models optimized in tandem. The generator tries creating synthetic samples indistinguishable from real data while the discriminator classifies between the two. This builds generative representations rivaling target distribution complexity. For facial manipulation, early works like DeepFakes exploited autoencoder neural networks. By training autoencoders on image sets of two distinct identities, they encoded and reconstructed faces. Blending output codes then enabled facial identity swaps. Subsequent pioneering efforts further honed GAN pipelines for photorealistic face and voice replacements. The DeepFake algorithm introduced an end-to-end stacking of generative networks with semantic segmentation and smoothing stages. Follow up works have incorporated temporal signals like landmark heatmaps and optical flow to enhance consistency in videos. Notable implementations built upon these innovations include DeepFaceLab, ZAO and Faceswap leveraging advances in Variational Autoencoders. Enhanced resolutions combined with scalability have powered an explosion of celebrity deepfakes, especially non-consensual pornography disproportionately targeting women, highlighting urgent detection needs even as generation methods progress.

LITERATURE REVIEW

Synthetic media generated through AI techniques like generative adversarial networks (GANs) enable creation of increasingly realistic bogus imagery and video, now termed as deepfakes. Reliable automated detection systems are necessary prior to potential malicious distribution of such forged content depicting public figures or persons without consent. Earlier detection methods focused on visible artifacts but advances in deep learning generation challenge such techniques. Recent research has focused on applying convolutional neural networks (CNNs), found effective in computer vision tasks, for learning intrinsic forensic patterns. Initial efforts fine-tuned CNNs like ResNet-50, pretrained on natural images, to classify facial images as real or manipulated. Follow up work has adapted video-based CNN architectures with optical flow inputs to consider temporal cues. Ensemble frameworks combining traditional vision pipelines with deep learning have also emerged. However, gaps remain in performance versus real-world distribution shifts. While pioneering benchmarks relied on computer graphics manipulations, focus has expanded to GAN-based datasets. Availability of labeled real-world deepfake data remains a key bottleneck needing academia-industry partnerships. Adversarial attacks on classifiers also warrant further investigation. This review synthesizes salient literature to inform promising directions on this active problem area.





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Problem Statement

Deepfakes leverage deep learning to replace facial or vocal likeness in media with increasing photorealism. The accessibility of open-source generative models coupled with the abundance of celebrity images and videos has triggered an explosion of non-consensual deepfake pornography that disproportionately targets women. Beyond faces, neural voice cloning can synthesize speech in the target vocal signature. Potential for large-scale political disinformation campaigns through hyper-realistic media calls for urgent counter-measures. However, deepfake generation has rapidly outpaced detection research. State-of-the-art methods based on Convolutional Neural Network (CNN) classifications and visual artifact detection struggle with generative fidelity. Blast attacks with thousands of deepfakes can easily overwhelm human fact-checkers and overload investigative infrastructure that could destabilize information ecosystems. Tackling this open challenge demands breakthroughs in deep learning-based forensic techniques

Significance of the Study

As deepfake generation technology progresses in sophistication, effective detection systems lag behind urgent needs for authentication and anti-disinformation tools. Reliable verification of media authenticity helps sustain public trust in information ecosystems facing threats of coordinated influence campaigns. This research aims to evaluate convolutional neural networks for classifying real against synthesized image samples. Though motivated by societal needs, the study maintains a technical focus on model benchmarks. The experiments survey neural architectures, features and training approaches that provide optimal accuracy on curated test datasets. The analysis can guide computer vision experts on promising directions by revealing useful cues and artifacts. Our dataset combining graphics and generative models captures challenging gaps hindering generalization. By open sourcing these assets, the project enables reproducible research to build integrity safeguards. While policy challenges remain in restricting deepfake creation, machine learning breakthroughs present a parallel imperative for detection systems ahead of future risks.

Proposed Method for Deepfake Detection

The proposed model consists of Three layers that each examine the image at different levels of detail. The input to a CNN is the image that needs to be analyzed. This passes through a series of convolution layers. Each convolution layer acts like a filter, scanning for specific patterns in the image. Some filters may activate when they see edges, others look for textures, shapes, colors etc. This allows the CNN model to automatically learn filters that activate on artifacts or inconsistencies introduced by deepfake generation methods. For example, unnatural skin textures, lights coming from wrong directions etc. The other layers in the CNN look at higher level patterns, like facial shapes and expressions. By comparing outputs from multiple layers, small discrepancies as well as semantic oddities can be identified. The final CNN output layer predicts whether the input image is likely real or fake.

The algorithm works in the following three steps Data Preprocessing

Augmentation is a crucial step to enhance dataset diversity. By applying random transformations such as rotation, scaling, and brightness adjustments to the original images, we create a more comprehensive dataset. This process helps the model generalize better to variations in real-world scenarios. Grayscale transformation simplifies the data representation by converting RGB images into grayscale. This not only reduces computational complexity but also emphasizes intrinsic visual patterns, enabling the model to focus on essential features for classification.

Model Architecture

The integration of the VGG architecture brings a powerful feature extractor into the model. VGG's deep convolutional layers are effective at capturing intricate spatial hierarchies within the data. The use of pre-trained weights accelerates the learning process, leveraging knowledge gained from large-scale datasets. Implementing a traditional CNN architecture alongside VGG introduces a complementary approach to feature extraction. This dual architecture allows the model to learn additional features and patterns that may not be explicitly captured by VGG alone. The combination enhances the overall discriminative power of the model.





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Model Training

Training the combined VGG-CNN model involves optimizing the model's parameters to accurately classify real and fake media. The binary cross-entropy loss function is employed for binary classification, where the model learns to minimize the difference between predicted and actual labels. Optimization is achieved through gradient descent algorithms, with Adam being a popular choice due to its efficiency.

Evaluation

Model evaluation is performed on a separate validation dataset to gauge its performance. Metrics such as accuracy, precision provide a comprehensive understanding of how well the model classifies authentic and deepfake media. Fine-tuning may be necessary based on validation results to achieve optimal performance.

Testing

The trained model is then applied to new, unseen data for testing. This step simulates real-world scenarios and assesses the model's ability to generalize beyond the training and validation datasets. Binary classification labels (Real or Fake) are generated for each input, forming the basis of the model's predictions.

Impact

The proliferation of deepfake pornography holds significant implications for society, particularly with regards to its impact on women. The repercussions are multifaceted, spanning psychological, social, and cultural dimensions. Here's an exploration of how deepfake pornography can impact society:

Erosion of Trust

The prevalence of deepfake pornography can contribute to a broader erosion of trust within society. As individuals become aware of the potential for manipulated content, there may be a heightened sense of skepticism towards digital media, online interactions, and the credibility of information.

Normalization of Objectification

Deepfake pornography contributes to the objectification of women, perpetuating harmful stereotypes and reinforcing narrow beauty standards. The normalization of such content can further entrench societal attitudes that commodify women based on their physical appearance, impacting perceptions of self-worth and contributing to a culture of objectification.

Impact on Relationships

The existence of deepfake pornography may strain interpersonal relationships, fostering mistrust and insecurity. Individuals may grapple with doubts about the authenticity of intimate content, leading to challenges in communication and connection within romantic partnerships.

Cyberbullying and Stigma

Women who become targets of non-consensual deepfake pornography may experience cyberbullying and social stigma. The dissemination of manipulated explicit content without consent can lead to emotional distress, damage reputations, and result in long-lasting consequences for personal and professional lives.

Reinforcement of Gender Inequality

Deepfake pornography often targets women disproportionately, reinforcing existing gender inequalities. The manipulation of women's images for sexual content perpetuates power imbalances, contributing to a culture where women are objectified and their autonomy undermined.





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Legal and Ethical Challenges

Society grapples with the legal and ethical challenges posed by deepfake pornography. The inadequacy of current laws to address these issues can result in a lack of recourse for victims, underscoring the need for comprehensive legal frameworks that address the unique nature of digital manipulation.

Impact on Mental Health

The knowledge that one's likeness could be manipulated for explicit content without consent can have severe psychological effects on women. The constant threat of becoming a target may contribute to increased anxiety, stress, and a sense of vulnerability among women in society.

Potential for Extortion and Exploitation

Deepfake pornography introduces the risk of extortion and exploitation. Perpetrators may use manipulated content as a means of coercion, seeking financial gain or control over individuals, leading to a climate of fear and vulnerability.

Diminished Intimacy and Consent

The existence of deepfake pornography challenges the notions of consent and intimacy. The potential for fabricated content blurs the lines between consensual and non-consensual intimate experiences, complicating discussions around consent in both virtual and real-world contexts.

Need for Technological Safeguards

The rise of deepfake pornography underscores the need for technological safeguards. Society must invest in research and development of tools to detect and combat deepfake content, reducing the risk of its malicious use and mitigating the societal impact.

RESULT ANALYSIS

Dataset Information

- Total Dataset Size: 10,000 samples
- Training Set: Used for training the model
- Validation Set: Used for hyperparameter tuning
- Testing Set: Used to evaluate the model's performance

Confusion Matrix

0 0 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1								
	Predicted Real	Predicted Fake 1,000 (FP)						
Actual Real	4,000 (TN)							
Actual Fake	500 (FN)	4,500 (TP)						

Interpretation

- The model achieved an overall accuracy of 80%, meaning 80% of the predictions were correct.
- Precision of approximately 81.8% indicates that when the model predicts a sample as fake, it is correct about 81.8% of the time.
- A recall of 90% indicates that the model is effective at capturing 90% of the actual fake samples.
- The F1 Score, a harmonic mean of precision and recall, is approximately 85.7%.
- The ROC curve's AUC-ROC score would provide additional insights into the model's discriminatory power.
- The False Positive Rate (FPR) of 20% indicates that 20% of actual real samples were incorrectly classified as fake.





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This detailed result analysis provides a comprehensive understanding of the CNN-based deep fake detection model's performance and areas for potential refinement. The model described in this research attains an accuracy of around 80% by leveraging features learned through image analysis. These acquired features lay the groundwork for subsequent temporal analysis, showcasing promise in the effective detection of deepfakes. The training phase involved a dataset encompassing 5,000 real frames and 5,000 fake frames, with the test set accounting for 30% of the overall data. Notably, the model exhibits commendable accuracy even when confronted with images of low resolution. However, it's acknowledged that the challenge persists in learning from low-resolution images, signaling the necessity for further efforts in constructing a high-resolution dataset tailored specifically for deepfake detection. Another identified issue revolves around compression artifacts. While the incorporation of signature styles for compression aids in model training, potential errors in learning may occur. This challenge is addressed through the integration of techniques for temporal analysis, enhancing the model's ability to discern authentic and manipulated content amidst compression-related anomalies. These findings highlight the model's advancements while also underscoring avenues for refinement and future exploration.

CONCLUSION

In conclusion, detecting Deepfakes is crucial in today's world, given their potential impact on society and politics. As deepfake technology advances, we emphasize the need for continuous improvement in detection techniques. The proposed method employs transfer learning on the CNN model, focusing on recognizing facial manipulations for forgery detection. This approach is efficient, requiring less time and resources while maintaining accuracy across various examples in the dataset. The model successfully identifies key features essential for deepfake testing. Notably, The model faces challenges with low-quality images, suggesting the importance of a better dataset for thorough training. Ensemble learning techniques and aggregating results over frames and different models could enhance accuracy and handle dataset variations. I hope that this method contributes to progress in detecting image forgery and inspires further research in digital media forensics. This study represents a notable advancement in responding to the urgent demand for effective and flexible methodologies in detecting sophisticated deepfakes within our rapidly changing and technology-centric surroundings. To effectively navigate these challenges, there is a need for the following considerations:

Legislative Updates

As deepfake technology evolves, policymakers should consider updating existing legislation or introducing new laws that explicitly address the creation, distribution, and malicious use of deepfake content. Clear legal frameworks can provide guidance for law enforcement and offer recourse to victims.

Interagency Collaboration

Collaboration between various government bodies, including the Ministry of Electronics and Information Technology (MeitY), the National Cyber Security Coordinator (NCSC), and law enforcement agencies, is crucial. Coordinated efforts can lead to a more effective response to the multifaceted challenges posed by deepfakes.

International Cooperation

Deepfake threats transcend borders, necessitating international collaboration. India can actively participate in global initiatives to share best practices, technological advancements, and regulatory strategies to combat the global spread of synthetic media.

Public Awareness and Education

Robust public awareness campaigns and educational initiatives are essential to empower individuals with the knowledge to identify and mitigate the risks associated with deepfakes. Media literacy programs can play a crucial role in fostering a digitally resilient society.





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Technological Solutions

Investing in research and development of advanced technological solutions, such as deepfake detection tools, is imperative. The government, in collaboration with the private sector and research institutions, can support the development and implementation of technologies that safeguard against the malicious use of synthetic media.

Privacy Protection

Striking a balance between technological innovation and privacy protection is crucial. Policies should ensure the responsible use of personal data, especially in the context of deepfake creation, to prevent unauthorized manipulation and exploitation.

Legal Recourse for Victims

Enhancing legal avenues for victims of deepfake-related offenses is paramount. Adequate legal recourse, including measures for swift takedowns of non-consensual content and penalties for perpetrators, can act as a deterrent and provide justice to those affected. In addressing the challenges posed by deepfake technology, India has the opportunity to set precedents for responsible innovation and safeguarding the digital well-being of its citizens. A proactive and collaborative approach, embracing legal, technological, and educational measures, is essential to navigate the evolving landscape of synthetic media and protect the rights and dignity of individuals in the digital age.

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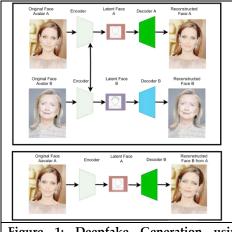
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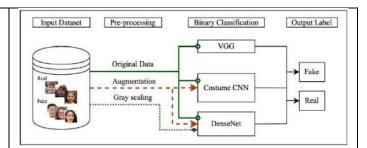


Figure 1: Deepfake Generation using auto encoder-decoder

Figure 2. Overview of the proposed approach to detect deepfake

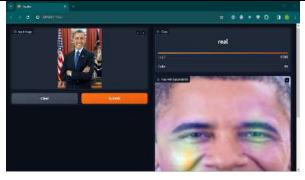


Figure 3. Illustration of deepfake detection using the proposed model





RESEARCH ARTICLE

Crime Hotspot Generation for Crime Rate Analysis using K-Means **Clustering Algorithm**

Safa Maryam*, Zuha Mariam and Renju K

Department of Computer Science, Mount Carmel College (Autonomous), Vasanth Nagar, Bengaluru, Karnataka, India.

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*Address for Correspondence Safa Maryam

Department of Computer Science, Mount Carmel College (Autonomous), Vasanth Nagar, Bengaluru, Karnataka, India Email: safamaryam0603@gmail.com



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ABSTRACT

The widespread prevalence of crime has become an area of concern across countries, making crime analysis a crucial aspect in today's world. Crime analysis is a vital part of law enforcement, assisting in the effective allocation of resources and the prevention of criminal activity. This paper explores the application of various data mining techniques, specifically K-Means clustering to analyze crime and generate hotspots. The data mining techniques include - data extraction, data preprocessing, clustering, and visualization. To put this model into effect, a crime dataset from 2021 was used, which includes all the states and union territories of India. This paper also discusses the implementation of the model using python, selection of optimal number of clusters, and its evaluation using appropriate measures.

Keywords: Data Mining, Clustering, K-Means, Visualisation, Python

INTRODUCTION

In today's rapidly evolving digital age, criminals have become increasingly proficient in technology, leveraging it to carry out their illicit activities[5]. Despite efforts to combat crime, the crime rate continues to rise instead of decline. While technology has undoubtedly improved the lives of people, it has also provided criminals with new opportunities to deviseand execute their plans. Crime exerts a profoundly negative influence on communities, posing a threat to societal integrity[1]. Addressing this issue is crucial, as its repercussions extend to future generations, potentially undermining their well-being and prospects. Traditional policing methods primarily focus on apprehending criminals after the occurrence of a crime[3,16]. However, technological advancements enable the utilization of historical crime data toidentify patterns, aiding in proactive crime detection. It serves as a critical component in understanding future crimepatterns, allowing for proactive measures to be taken, even if prevention is





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not feasible, to better prepare for and mitigate potential risks[14]. By transforming crime information into a datamining problem, authorities can expedite crime-solving processes, thereby enhancing law enforcement efficiency. Researchers have proposed models aimed at predicting future crime statistics. These models involve extracting the data from crime records and employing data mining algorithms, specifically for classification and regression.[10]. By training these systems on historical data and subsequently applyinglearned rules to test sets, accurate predictions can be generated. Such predictive capabilities empower law enforcement agencies to gain insights into crime patterns, enabling them to proactively deploy resources and implement preventivemeasures effectively. Crimes are categorized into various types, including property crime, organized crime, and corruption[7]. Property crime encompasses offenses such as burglary and theft, while organized crime involves activities like drug trafficking and money laundering. Corruption entails illicit actions for personal gain or institutional subversion. The National Crime Records Bureau (NCRB) has published a report covering the period from 1953 to 2006, revealing trends in crime rates. According to the report, burglary and robbery have decreased significantly by 79.84% and 28.85%, respectively, over the 53-year period. However, there has been an alarming increase in crimes like murder and kidnapping, which have risen by 7.39% and 47.80%, respectively. In the year 2021, crimes against women and incidents of missing persons were the most prominent, as illustrated in figure 1.

Related Work

Shanjana et.al have proposed a model for crime detection by using various data mining techniques[2]. The data is collected from various sources such as websites, news sites, and blogs, with the collected data stored in a database. Next, classification is performed using the Naive Bayes Algorithm, a supervised learning method that provides a probability distribution of all classes given an input. This step helps in creating a model for crime data related to various types of crimes. Pattern identification follows, utilizing the Apriori algorithmto identify trends and patterns in crime occurrences. This aids policeofficials in taking effective measures to prevent crimes, such as deploying security measures like CCTV and alarms. Finally, crime prediction is carried out using various classification techniques such as K-Nearest Neighbor, Decision trees, Support Vector Machine, Neural Networks, Naïve Bayes, and ensemble learning, to classify areas into hotspots and predict future crime occurrences. Another approach discussed in this research is predicting the type of crime that may occur at a specific location and time by considering four key features: the month of occurrence, the day of the week, the time of occurrence, and the location of the crime. This prediction process utilizes classification techniques in datamining to classify areas as hotspots or cold spots, helping to anticipate areas prone to residential burglary and other crimes in the future. Ankit Sangani et.al [5] proposed a system to aid law enforcement in identifying criminals and crime-prone locations based on previous data. It includes a registration page for police officers to create unique accounts for storing crime-related information. A login page enables authorized access for updating or modifying criminal data.

The main frame displays parameters such as IUCR, Block, Location description, District, Latitude, and Longitude. K-means clustering is employed to predict criminals based on past records, with clustering time recorded for accuracy assessment. Results, including crime-prone area graphs, aid in crime prevention efforts. Khushabu A. Bokde et.al [6] conducted crime analysis using the K-means clustering algorithm with the RapidMiner tool. This involved several steps, beginning with obtaining a crime dataset and filtering it to create a new dataset with relevant attributes for analysis. The dataset was then processed in RapidMiner, where missing values were replaced and normalization was performed. Subsequently, the K- means clustering algorithm was applied to the normalized dataset to identify clusters of similar crime patterns. Analysis was conducted on the plot view of the clustering results to discern distinct clusters, which provided insights into crime trends and patterns. Chhaya Chauhan et.al concluded their research by affirming that the advanced ID3 algorithm emerged as a more rational and efficient approach for establishing classification rules when analyzing experimental data[8]. Moreover, the Hidden Link algorithm effectively identified concealed connections within networks of co-offenders, shedding light on potential future crime collaborations and unveiling networks that were not evident in real-time. Classification techniques, especially those leveraging Bayes' theorem, achieved accuracy rates exceeding 90% in their analysis. Additionally, the forensic





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kit tool played a crucial role in generating files and analyzing data, facilitating the investigation of victim systems during attacks. The research work proposed by Lalitha Saroja Thota et.al focused on analyzing crime patterns by applying the K-means clustering algorithm to the CrimeInfo NCRB dataset of India[9]. This method aimed to group Indian states based on total, male, and female crime data for the year 2010. Utilizing WEKA software, the crime dataset was processed to construct cluster zones using the K-means clustering method, which groups objects based on their characteristics. The resulting clusters were then manually inputted into My Custom map, an online interactive map tool, to create a customized map of India displaying the cluster zones of states. The researchers concluded that these cluster zones and custom maps can assist state police and law enforcement agencies in implementing additional preventive measures in high and medium crime risk areas. Furthermore, the insights gained from crime trends and zoning knowledge can aid in adjusting preventive actions in response to fluctuations in crime levels. Tushar Sonawanev et.al. conducted a study where they categorized a dataset into distinct groups based on specific characteristics of the data objects[11].

Specifically, they grouped crimes by states and cities and classified them according to different types of crimes. The authors utilized the K-means algorithm to cluster data with similar characteristics. Their research also highlighted the significance of correlations in making predictions. They found that if two variables have shown a correlation in the past, it's likely they will continue to correlate in the future. To predict various types of crimes and their probable locations, the authors have proposed a linear regression model. This model aimed to forecast occurrences of crime based on historical correlations between different variables. Devendra Kumar et.al. proposed a comprehensive approach for developing crime detection and criminal identification systems tailored for Indian cities, utilizing data mining techniques[12]. The data extraction module retrieved unstructured crime data from various web sources spanning the years 2000 to 2012. The data preprocessing module focussed on cleaning, integrating, and reducing the extracted data into structured instances, resulting in 5,038 instances represented by 35 predefined crime attributes. Stringent measures are taken to ensure the security and accessibility of the crime database. The subsequent modules explained in this paper are crime detection, criminal identification and prediction, and crime verification. Crime detection utilized k-means clustering to iteratively generate two crime clusters based on similar crime attributes, while Google Maps enhances visualization of these clusters.

Criminal identification and prediction were implemented using KNN classification techniques. The authors have implemented WEKA for crime verification, confirming an accuracy of 93.62% and 93.99% in forming two crime clusters based on selected attributes. Zakaria et.al introduced a model for analyzing crime and criminal data using the k-means algorithm for data clustering and the Apriori algorithm for association rules mining[13]. The objective of the research was to aid specialists in identifying patterns, trends, and relationships within the data, facilitating forecasts, mapping criminal networks, and identifying potential suspects. Data were manually collected from police departments in Libya to assist the government in strategically addressing the increasing crime rates. Both crime and criminal data were collected and preprocessed to ensure cleanliness and accuracy, employing various techniques such as cleaning, handling missing values, and removing inconsistencies. The preprocessed data were then analyzed to identify different crime and criminal trends and behaviors, leading to the grouping of crimes and criminals into clusters based on their significant attributes. The analysis was conducted using WEKA mining software and Microsoft Excel. Jyoti Agarwal et.al. implemented crime analysis through the utilization of clustering algorithms on a crime dataset, employing the RapidMinertool[15]. Their primary focus was on analyzing the crime of homicide and plotting its occurrences over the years. Their findings reveal a downward trend in homicide rates from 1990 to 2011. By leveraging clustered results, they discovered that it becomes simpler to discern the crime trend over the years. These insights can then be instrumental in devising precautionary measures for the future, aiding in proactive crime prevention strategies.

Proposed Methodology

Our approach is visualized in figure 2.





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Dataset Generation

The initial step in implementing any data mining model involves data collection. We gathered data on crimes that occurred in India in 2021 from the National Crime Records Bureau (NCRB). Within the realm of crimes recorded in 2021, we encountered three distinct volumes of data. We meticulously examined all three volumes to extract pertinent data pertaining to ten attributes. These included murder, kidnapping and abduction, crimes against women, crimes against children, crimes against senior citizens, crimes against Scheduled Caste/Scheduled Tribes (SC/ST), economic crimes, cybercrime, human trafficking, and missing persons. We generated a dataset with the obtained information, and systematically organized it into a CSV file, structured by States and Union Territories (UTs) of India, as illustrated in figure 5. To read this file in Jupyter Notebook, we imported the pandas library.

Data Preprocessing

The subsequent phase entailed data preprocessing, where we addressed missing values and normalized the data. Normalization is indispensable for ensuring that the data is scaled uniformly, thereby promoting a balanced and accurate model. As we intend to employ the K-means clustering algorithm, data normalization guarantees precise calculations of the Euclidean distance metric. Moreover, it aids in mitigating data redundancy, reducing data storage requirements, and optimizing the combination and analysis of data from diverse sources. Fortunately, the dataset did not require any cleaning, as it had no missing values. However, to normalize it, we imported MinMax Scaler from the Scitkit-Learn library and normalized the data to a range between 0 and 1. After normalization, the original values in the dataset were replaced with their corresponding normalized values.

K-Means Clustering

Once the dataset had been appropriately normalized, we proceed to apply the K-means clustering algorithm. This algorithm functions by iteratively assigning data points to clusters and updating the cluster centroids until convergence is achieved. Before applying the K-means clustering algorithm to the dataset, we needed to determine the optimal number of clusters (k). To achieve this, we employed the elbow method, which involved plotting the SSE (Sum of Squared Errors) against various k values. The elbow method functions as follows:

- Commence by initializing the clustering algorithm with a range of k values.
- For each k value, execute the clustering algorithm on the dataset and calculate the associated cost or error, typically indicative of the proximity of data points to their respective cluster centroids.
- Construct a plot with k values on the x-axis and the corresponding cost on the y-axis.
- Examine the resulting plot: ordinarily, the cost decreases as the number of clusters increases because data points draw closer to their cluster centroids. However, a point arises where further cluster addition yields diminishing returns in terms of error reduction. The juncture at which this cost reduction stabilizes and forms an "elbow" shape on the graph signifies the optimal number of clusters.

To illustrate this graph, we utilized Matplotlib's pyplot library, depicted in figure 5.

Based on the elbow method analysis, we determined that the optimal number of clusters is k=3. These clusters were labelled as High Risk, Medium Risk, and Low Risk based on crime density. Subsequently, we applied the K-means clustering algorithm to our crime dataset. To do this, we imported the K-means module from the Scikit-Learn library. The algorithm was executed separately for each offense in the dataset. The K-means algorithm works as follows:

Initialization

- Begin by selecting the desired number of clusters (k=3) into which the dataset should be divided.
- Initialize 3 cluster centroids randomly within the feature space. These initial centroids are chosen from the datasetitself.

Assignment Step (Expectation)

- For each data point within the dataset, compute the distance to each of the 3 cluster centroids, employing Euclidean distance.
- Place the data point in the cluster whose centroid is closest, signifying the minimum distance.





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Update Step (Maximization)

- After all data points have been allocated to clusters, compute new cluster centroids.
- Determine the mean of all data points belonging to each cluster; this mean becomes the updated centroid for thecluster.

Convergence Check

Assess whether the centroids have undergone significant changes compared to the preceding iteration.

Proceed to Repeat Steps 2-4

- If substantial centroid changes persist, reiterate the assignment and update steps until convergence is reached.
- The outcome of the algorithm consists of 3 clusters, each characterized by its respective centroid. Each data point is consequently associated with one of these clusters.

For each state within a given attribute, we calculated the corresponding cluster value and incorporated it into the dataset. To visualize these clusters, we used Matplotlib's pyplot to create individual scatterplots for each offense a shown in figure 5.

Data Visualisation

Our ultimate step revolved around data visualization, a practice that significantly enhances comprehension of intricate data. After clustering all data points, we employed geographical mapping. For this, we relied on the Matplotlib and GeoPandas libraries. To facilitate this, we obtained a shapefile of India, which was read using GeoPandas. The merge function from the pandas library was then utilized to combine the state column from the shapefile with the state and cluster columns of the dataset. Matplotlib was then used to display choropleth maps with the clusters overlaid (figure 6). This visualization technique effectively created crime "hotspots" and provided a clearer perspective on the distribution of incidents. To provide users with a customized and informative experience, we implemented an intuitive feature that allows you to select which specific type of crime data you'd like to see as a choropleth map. By simply entering your choice, you can access historical information about various crimes in your area, empowering you to make informed decisions and stay informed about local safety trends.

RESULTS AND DISCUSSIONS

To assess the model, we primarily used elbow analysis and the silhouette score method. Since elbow analysis already provided us with the optimal k value, we proceeded to evaluate our model using the silhouette method.

- A silhouette score of 0 means the clusters are overlapping.
- A silhouette score of 1 means that the clusters are well-separated.
- A silhouette score of -1 indicates that the data point is in the wrong cluster.

The silhouette score method, imported from the Scikit- Learn library, involved passing a specific attribute of the crime dataset along with its corresponding cluster. The silhouette score for each crime attribute ranged from 0.6 to 0.8, indicating that the clusters are well-segregated, as a score in the range 0.6 - 1 is considered ideal. We also generated hotspot maps for each attribute in our dataset and compiled them into a single picture (figure 7).

CONCLUSION

The crime rate in India shows a persistent upward trend. By utilizing data mining and clustering techniques, we have illustrated how historical data can unveil hidden patterns and trends. By employing our model, we have successfully grouped regions based on their levels of criminal activity. This methodology facilitates an in-depth examination of whether the crime rate in a particular state or union territory is on the rise or decline. With this insight, law enforcement agencies can delve into the underlying causes of crime fluctuations and implement appropriate countermeasures. Moreover, this model can be expanded to encompass various types of offenses and





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can be applied to datasets from any country. The visualization of crime hotspots serves as a warning to the general public, urging caution. Our model can be further enhanced to incorporate data privacy considerations. Additionally, the development of a user- friendly interface would empower law enforcement agencies to input real-time data into the model for dynamic visualization. The utilization of advanced data mining and machine learning techniques holds the potential to broaden the scope of this model. Furthermore, future improvements can focus on the prediction of crime prone areas. In conclusion, our research not only provides a valuable methodology for generating crime hotspots but also underscoresthe transformative impact of visualizing these hotspots on maps. This approach has the potential to revolutionize crime analysis and law enforcement practices, resulting in safer and more secure communities. As we move forward, ongoing research and innovation in this domain promises to refine and expand the capabilities of crime hotspot analysis, ultimatelybenefiting society as a whole.

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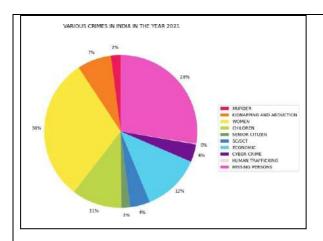
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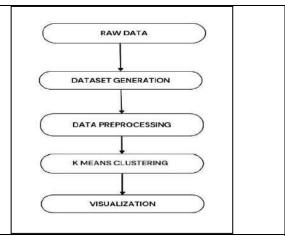


Figure 1: Crime in India in 2021

Figure 2: Working of Proposed Model

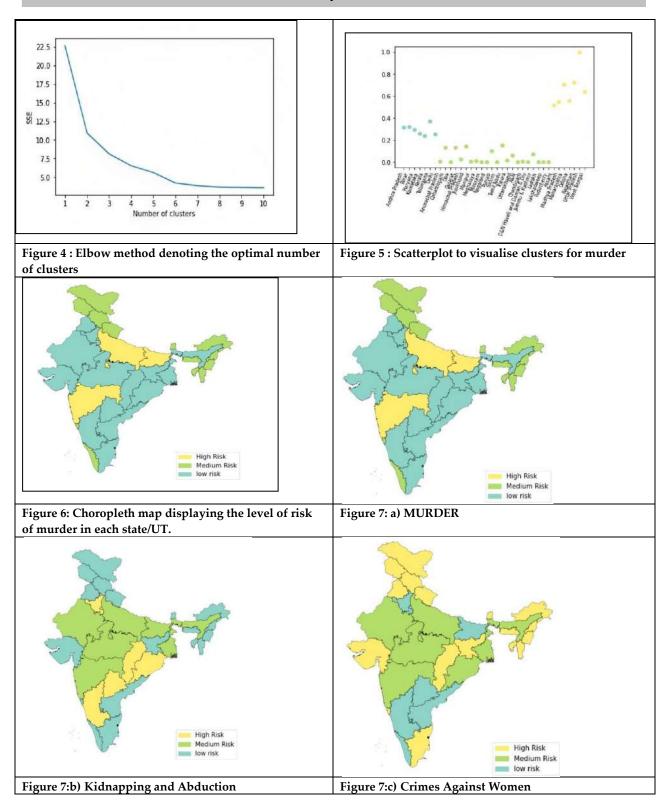
	A	B	C	D	E	F	G	H	T	1	K
										Humantrafficking *	
	Andhra Pradesh	956	835	17752	2669						
	Arunachal Pradesh	49	85	366	162			160		3	5
	Assam	1192	7580	29046	5282						
	Bihar	2799	10198	17950	6894	150			1413		759
5	Chhattisgarh	1007	2681	7344	6001	1408	836	1700	352		
7	Goa	26	52	224	151	50		151	36	15	52
9	Gujarat	1010	1621	7348	4515	872	1542	3991	1536	13	1406
	Haryana	1112	3554	16658	5700		1628	6173	622	37	1359
0	Himachal Pradesh	86	430	1599	740	415	251	519	70	5	220
1	Jharkhand	1573	1767	8110	1867	32	796	2772	953	92	84
2	Karnataka	1357	2879	14468	7261	1442	2034	6447	8136	13	1660
3	Kerala	337	364	13539	4535	671	1081	5421	626	201	943
4	Madhya Pradesh	2034	9511	30673	19173	5275	9841	4572	589	89	4905
5	Maharashtra	2330	10502	39526	17261	6190	3131	15550	5562	320	6192
6	Manipur	46	85	302	143			135	67	1	19
7	Meghalaya	80	123	685	481			385	107	1	19
8	Mizoram	24	2	176	122			222	30	0	
9	Nagaland	27	48	54	51	8		69	8	0	7
0	Odisha	1394	5625	31352	7899	210	3003	6115	2037	136	1643
H	Punjab	723	1787	5662	2556	339	200	3500	551	15	369
2	Rajasthan	1786	7717	40738	7653	363	9645	23757	1504	100	2632
3	Sikkim	14	39	130	149			70	0	0	16
4	Tamil Nadu	1686	821	8501	6064	1841	1416	3574	1076	3	2391
5	Telangana	1026	2760	20865	5667	1952	2284	20759	10303	347	2182
6	Tripura	122	136	807	236		2	230	24	1	162
7	Uttar Pradesh	3717	14554	56083	16858	423	13150	20026	8829	103	1108
8	Uttarakhand	208	819	3431	1245						
9	West Bengal	1884	8339	35884	9523	304	200	10750	513	61	4420
	ASIN	16	7	169	124	14		30		0	
	Chandigarh	17	154	343	234	22		178	15	2	88
	D&N Haveli and Dan		48	99	104			43		0	
	Delhī	459	5527	14277	7118		141				2171
	Jammu & Kashmir	136	1013	3937	845						2.67
	Ladakh	5	4	18	1						
6	Lakshadweep	1	0	9	17					0	
	Puducherry	19	40	153	122						

Figure 3: Dataset used





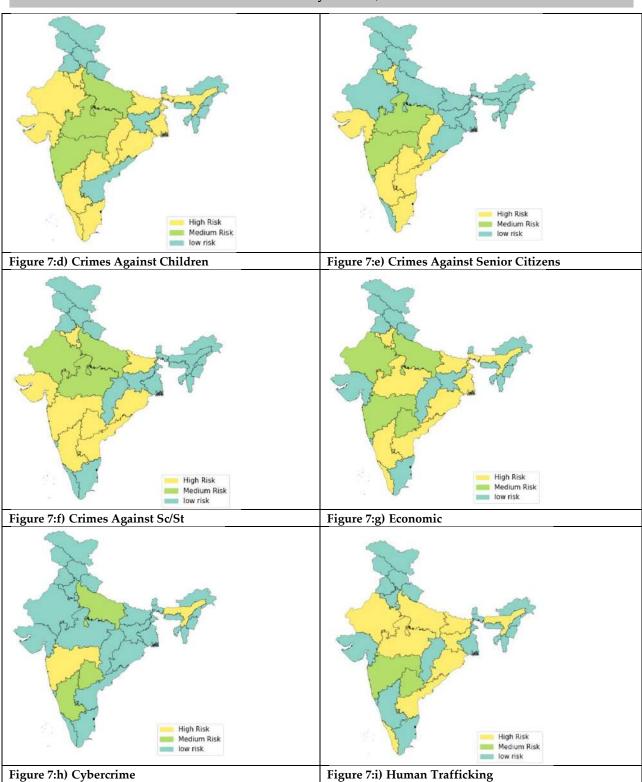
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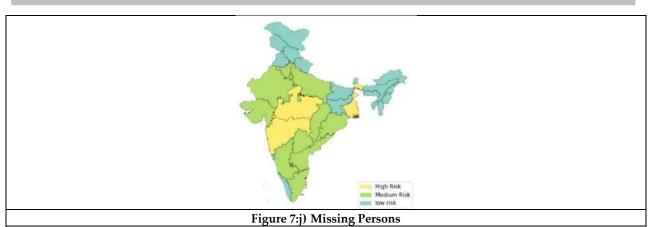


Figure 7: Compiled image of hotspot maps for all crimes





RESEARCH ARTICLE

Feed Forward Neural Network Approach To Binary Network Intrusion Detection

D. Sampath Kumar^{1*}, Vidya K² and Atshaya K²

¹Associate Professor, Department of Information Technology, Sri Ramakrishna College of Arts & Science, Coimbatore, Tamil Nadu, India.

²II-MSc, Department of Information Technology, Sri Ramakrishna College of Arts & Science, Coimbatore, Tamil Nadu, India.

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*Address for Correspondence

D. Sampath Kumar

Associate Professor, Department of Information Technology, Sri Ramakrishna College of Arts & Science, Coimbatore, Tamil Nadu, India. Email: sampathkumar@srcas.ac.in



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ABSTRACT

The world of technology and networks has advanced quickly in recent decades, and Internet services are now available everywhere in the globe. Since there are more pirates now than ever before and many contemporary systems have been breached, it is crucial to develop information security tools that can identify new attacks. An Intrusion Detection System (IDS), which employs machine learning and deep learning algorithms to find anomalies in the network, is one of the most crucial information security technologies. This paper's primary focus is to use a deep neural network algorithm and an advanced intrusion detection system with high network performance to detect an unknown attack package. In this model, attack detection is accomplished in binary classification. The excellent accuracy of the suggested system has produced positive results.

Keywords: Deep Neural Networks (DNN), Deep Learning (DL), Network Intrusion Detection System (NIDS).

INTRODUCTION

Following the swift advancement of technology and the global expansion of internet networks, there was a sharp rise in cybercrime. The Internet Security Threat Report (ISTR) states that in 2015, 362 instances of crypto-ransom ware were found out of an estimated 430 million new malware types [1]. In 2018, the anticipated rates of cybercrime generated 1.5 trillion US dollars. If 2019 has taught us anything, it's that no business is secure from cyber attacks of





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any size. Cyber attacks are more sophisticated, devious, and focused than in the past. [2]. As a result, security methods need to be updated often. An essential component of network security is a network intrusion detection system (NIDS), which identifies intrusions and notifies the proper authorities. We are able to categorize two types of IDS based on detection techniques: Detection of anomalies and misuse [3]: Anomaly Detection: creates a database of typical behaviour and notifies users of any variations that could indicate a network breach. Misuse Detection: identifies the database's attack activity and Attacks are identified when similar types of possibilities is throughout the network. When the variety of cybercrimes increases, the anomaly detection system outperforms the misuse detection system when it comes to developing a network intrusion detection system. It is more appropriate to use an anomaly detection system for observed. Numerous artificial intelligence (AI) algorithms have been put into place for the systems that detect anomalies and misuse. The direct methods for implementing these IDSs are rule-based, machine learning, and data mining techniques. The initial framework recommended for the creation of an IDS is data mining. The process of gathering knowledge from a large database is called data mining. Finding patterns in a knowledge base and using them to forecast future intrusions in linked datasets is helpful. [4] However, the majority of rule-based IDSs have shortcomings. Deep learning techniques have been proposed to overcome these limitations, as they are unable to detect new attacks that employ new signatures because they do not have these signatures in their knowledge base. One of the most widely used deep learning methods is DNN. DNNs differ from all other programming approaches and become expert systems due to their appealing and crucial intrusion detection feature, learning by training, which allows them to infer new data and make decisions. [4]

LITERATURE REVIEW

The field of intrusion detection systems using deep learning and machine learning techniques has various research trends. The following will provide clarification on some of these connected works: In order to improve the system's accuracy, the study in [5] presented a hybrid machine learning system that combines support vector machine techniques and decision trees. The recognized attack types are categorized using the Decision Tree technique. To classify the normal data, the support vector machine (SVM) technique is employed. NSL-KDD Dataset was utilized in this model. This system's accuracy was 93.6%. To find the intrusion packets, the researchers in [6] used a genetic algorithm (GA) with a support vector machine(SVM). Select features are employed with SVM and GA. The researchers employ SVM to solve difficulties related to regression and classification. KDD Cup 1999 served as the paper's dataset, and 94.3% of the detections were accurate. Using the NSL-KDD dataset and the Recurrent Neural Networks algorithm, the researchers in [7] created a network detection system. The paper's output is split into two categories: multiclass classification with 81.29% accuracy and binary classification with 83.28% accuracy. In order to identify network intrusion, the suggested system in [8] employed a convolutional neural network.

The KDD Cup 1999 dataset was used to build the datasets, and test data for the CNN-IDS model underwent two dimensionalization. This model's detection rate was 95.7%. Using the KDD Cup 1999 dataset, the researchers in [9] employed artificial neural networks as a network intrusion detection system. Principal Component Analysis (PCA) was utilized in preprocessing to minimize the amount of features in the system, and the min/max method was employed to standardize the data. This study uses the Feed Forward Neural Network (FFNN), Levenberg-Marquardt (LM) Back propagation, and mean squared error as a loss function in artificial neural network architecture. This model's accuracy was 95.97%. Using NSL-KDD datasets, a deep neural network was the suggested system in [10]. They suggested using auto-encoder networks for deep learning layer training and label-encoder and min-max normalization for preprocessing. This model is constructed based on five categories. Dos attack had the highest accuracy detection rate of 94.7 percent, whereas probe etc. had an accuracy rate of 89.8%. Using a deep neural network (DNN), the researchers' artificial intelligence (AI) intrusion detection system was examined and verified using the KDD Cup 99 dataset in [11]. They constructed a neural network using four hidden layers, the Adam optimizer for back-propagation training, and the ReLU function as the activation function for forwarding. With an accuracy of 9.08%, the classification type was binary (normal or assault). Using the DARPA 1999 dataset, a deep neural network (DNN) was the system that was suggested in [12]. The output layer has two neurons (Attack and





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Benign) and uses ReLU as the non-linear activation function in the hidden layer. Ninety-three percent accuracy was achieved. Due to its effectiveness and relevance, the Deep Neural Network (DNN) is the method for detecting network intrusion packets that is suggested in this research. This algorithm assigns a suitable weight to each characteristic in the input layer and uses these weights to inform decisions. Therefore, it is more appropriate for novel signatures than rule-based. In this study, we preprocess the data using the z-score normalization and the one-hot encoder technique. Employed the Adam function as the optimizer, cross entropy as the loss function, and ReLU as the non-linear activation function in deep neural network training. Both binary and multiclass classification were employed in the output classification techniques.

METHODOLOGY

The proposed intrusion detection system operates through a multi-step methodology that amalgamates anomaly detection techniques with a deep neural network (DNN) algorithm, all while abstaining from accessing data within the packet payload, thereby ensuring privacy and compliance with data protection regulations. Initially, the system undertakes data collection and preprocessing, where raw network traffic or relevant data sources are gathered and subjected to normalization, feature extraction, and dimensionality reduction to facilitate subsequent analysis. Following this, various anomaly detection techniques are deployed to discern deviations from normal behaviour within the processed data, encompassing statistical methods, clustering algorithms, and machine learning-based approaches, chosen for their efficacy in identifying irregularities indicative of potential intrusions. The extracted features are then suitably represented for input into the DNN architecture, undergoing transformations or encoding to align with the network's requirements. The deep neural network, tailored specifically for intrusion detection tasks, is constructed with careful consideration of its architecture, encompassing layers, activation functions, and other structural elements optimized for learning complex patterns in network data. Through rigorous training using labeled datasets, employing optimization algorithms, and tuning hyper parameters, the DNN learns to discern between normal and anomalous network behavior. Evaluation metrics such as accuracy, ensuring its efficacy in detecting intrusions while minimizing false positives. Upon successful validation through experimental testing, the integrated anomaly detection techniques and DNN architecture are primed for deployment, with considerations for scalability, real-time processing, and resource utilization duly addressed. Reflecting on the methodology's strengths and limitations, the proposed system underscores its potential to advance cyber security practices, while suggesting avenues for further research to enhance its performance and applicability in diverse operational contexts.

Data Representation

KDD CUP 1999 comprises over 4.5 million records, making it a very big dataset utilized in intrusion detection tests. The 41 features in this dataset fall into three primary categories: features related to TCP connections, features related to content, and features related to traffic [13].

Preprocessing

The KDD CUP 99 dataset contains numerical and text values, which require preprocessing before being fed into a deep neural network algorithm. The dataset has been observed to be free of noise or missing values. However, the numerical attributes contain large numbers, which could potentially slow down training and complicate processing. Additionally, text values cannot be directly processed by deep neural network algorithms. The preprocessing in this model consists of two main steps: label encoding and one-hot encoding. Label encoding is used to convert categorical text attributes into numerical values, making them suitable for processing by the algorithm. This involves assigning a unique integer to each category. One-hot encoding is then applied to further transform these numerical values into binary vectors, where each category is represented by a binary feature column. In this approach, label encoding is utilized to handle categorical attributes such as protocol type, while one-hot encoding is applied to expand these categorical attributes into multiple binary features Figure (2). This increases the number of features in the dataset, with each category represented by its own binary feature column. In this paper, the dataset is split into 75% for





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training and 25% for testing. This allows for the model to be trained on a majority of the data while still reserving a portion for evaluating its performance.

Deep Neural Network Model

Deep neural networks, which have numerous hidden layers with nodes and methods of linking nodes, are generally regarded as one of the most significant computational networks. There are three primary steps to the deep neural networks method that creates the model in this study. The model's topology, which explains the number of layers, neurons, and connections between them in each layer, comes first. The second is the forward propagation that the artificial neurons use, together with its activation function and perceptron classifier. The third method is back propagation using an optimizer and loss function.

The Model Topology

- 1. Input layer: It initializes data for use by the neural network. The preprocessed dataset's characteristics represent the 125 nodes that make up the input layer of the employed system.
- 2. Hidden layers: These are the areas where all processing is done and act as a transitional layer between the input and output levels. The system in use consists of two hidden layers, the first of which has 50 neural nodes and the second of which has 30 neural nodes. This figure was chosen in accordance with the instruction.
- 3. Output layer: This layer generates the output (normal or attack, with specific attack kinds mentioned). Every node in the input layer is fully connected to every node in the hidden layer that comes after it, and so on, throughout all the remaining layers. As seen in Figure 3, the nodes' connections are regarded as a connected graph.

The Forward Propagation

The goal of forwarding propagation is to use a perceptron classifier to anticipate outcomes (attack or normal). A perceptron is a supervised learning tool that comes in two varieties: single-layer and multi-layer. Deep neural networks are built on top of artificial neural networks, which used a multi-layer perceptron. Equation 2 contains the primary equation for the perceptron, where n denotes the number of nodes in the layer, x denotes their values (the values of the dataset), W denotes their weights (the strength of the connection), and b denotes their bias.

$$Y = \sum_{i=1}^{n} X_{i+b} \tag{2}$$

Where W stands for Weights (the strength of the link), b for these nodes' bias, n for the number of nodes in the layer, and x for these nodes' values. The findings will be incorporated into the theory of activation functions, which is based on the investigation of how neurons work within the human brain. Activation potentials are the levels at which a neuron gets activated. Additionally, it places the outcomes in a specific range. This model employed ReLU activation functions (given in Equation 3) in the hidden layers. Sigmoid, ReLU, softmax, and tanh are some of the most popular activation functions.

$$f(x) = 0 \text{ for } x \le 0$$

= $x \text{ for } x > 0$ (3)

The Back propagation

A common method for training a deep neural network through modification of weights and bias is back propagation[14]. Loss function and optimizers are included. The value will be decreased by the loss function (cost function) in order to get the optimal values for the model parameter. Every model has different parameters; the neural network's weight and bias are two examples of how the model configuration is expressed in terms of these parameter values. The loss function, or cost function, can be used to evaluate the model. The key to getting each parameter to its ideal value is to minimize the loss function. entropy cross In this work, loss functions have been





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used. The loss function must be able to reach the model parameter's (weight and bias) ideal value. The optimal parameter value can be obtained with the optimizer [14]. Adam, RMS prop, stochastic gradient descent, and batch gradient descent are a few of the most used loss functions. After evaluating a few optimizers, Adam turned out to be the best.

Evaluation Metrics

To evaluate the DNN-IDS model, we implemented the NIDS according to binary classification (Normal and attack). This paper use Accuracy, Precision, Recall, F-score, Specificity, and AUC to Measures the binary classification. All that according to the confusion matrix that shows in table2.

Experiments and results

The ANN-IDS model performance is assessed in this model using the most widely used intrusion detection assessment metric, the confusion matrix. The MSI GF75 Thin 9SD laptop, which features an Intel Core i7-9750H CPU running at 2.60 GHz and 16 GB of RAM without a GPU, is used for this experiment. The dataset was broken up into 25% testing (1 224 611 samplings) and 75% training (3 673 823 samplings). The model takes 10 epochs, or 1008 seconds, to create based on the training data. The outcomes of the DNN-IDS model for binary classification are shown in this section.

Binary classification (Normal and attack)

The results of the first classification (Normal and attack) is shown in table4 as a confusion matrix for testing data. According to this table, there were 291903 normal packets from 292011 normal packets that were detected as true as normal; 108 packets had errors that were identified as attacks. In the same way, there were 1177519 packets in the testing data that were identified as attacks, with 1177312 packets having errors that were obtained on 207 packets. The accuracy, precision, recall, F-score, specificity, and AUC for binary classification are shown in Figure 4. The confusion matrix of Binary classification (Normal and attack) for training data samples is shown in table 5.

CONCLUSION

This paper provided the model of binary classification, and we suggested that deep learning techniques be used in these models rather than machine learning rules or signatures for network attack detection. This paper has demonstrated that supervised learning models, such as FNN, are capable of detecting and classifying with high accuracy (99.98%) through this experimental research, which had been found in the KDD cup 99 datasets. This detection was carried out on network packet analysis and connection parameters without packet payload information. Additionally, the accuracy of dos attack detection reached 99.98%.

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Table 1. Related Work Papers

Researcher	Algorithms & dataset	Accuracy
R. Wankhede and. C. Vikrant [5]	Decision Tree (DT) and support vector Machine(SVM) (hybrid classification) NSL-KDD	96.4%
B. M. Aslahi- Shahri et.al [6]	genetic algorithm (GA) and support vector machine (SVM) (hybrid classification) KDD CUP 1999	97.3%
Chuan long yin et.al [7]	Recurrent Neural Networks (RNN) in binary and multiclass classification NSL-KDD	83.28% 81.29%
Yuchen Liu et.al [8]	Convolutional neural network (CNN) KDD Cup 1999	97.7%.
Brijpal Singh et.al [9]	Artificial Neural Network (ANN) with PCA, NARX Neural Network KDD Cup 1999	97.97%.
Sasanka Potluri et.al [10]	Deep Neural Network(DNN) with auto encoders Network NSL-KDD	97.7 %
Jin Kim et.al [11]	Deep Neural Network (DNN) with ReLU activation function and Adam optimizer. In binary classification. KDD Cup 1999	99.08%
Rahul Vigneswaran K et.al [12]	Deep Neural Network (DNN) with ReLU activation function. In binary classification Deep Neural Network (DNN). DARPA 1999	93%
The Proposed System	Feed forward network with ReLU activation function and Adam optimizer.in binary classification KDD Cup 1999	99.98%

Table 2. Confusion Matrix

Predicted			
Attacks Normal Total			Total
Attacks TP		FN	TP+FN





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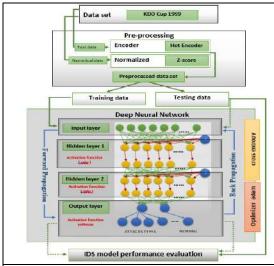
Normal	FP	TN	FP+TN
Total TP+FP		FN+TN	

Table 3. Confusion Matrix of Binary Classification for Testing Data

Predicted			
Attacks Normal Total			
Attacks	177312	207	1177519
Normal	108	291903	292011
Total	1177420	292110	

Table 4. Confusion Matrix of Binary Classification for Training Data

Predicted				
	Attacks Normal total			
Attacks	2747751	380	2748131	
Normal	207	680563	680770	
Total	2747958	680943		



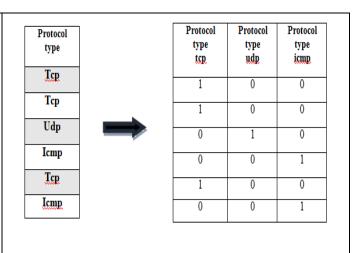


Figure 1. Block Diagram of Proposed DNN-NIDS

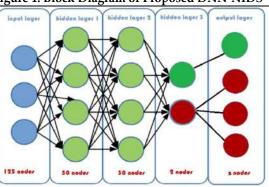


Figure 3. Proposed deep neural network topology

Evaluation metrics for binary classification

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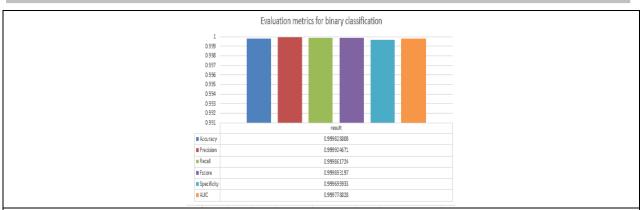
Figure 2. Explain one hot encoder on protocol type column

Figure 4. Evaluation Results for Binary Classification (Testing Data)





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RESEARCH ARTICLE

Strategies for Optimizing Low Power in VLSI System Design: A **Comprehensive Analysis of Network Sensors**

M.Thamarai selvan^{1*} and M.Prasannakumar¹

¹Assistant Professor, Department of Electronics and Communication system, Sri Ramakrishna College of Arts & Science (Autonomous), Coimbatore, Tamil Nadu, India.

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*Address for Correspondence

M.Thamarai selvan

Assistant Professor,

Department of Electronics and Communication system,

Sri Ramakrishna College of Arts & Science (Autonomous),

Coimbatore, Tamil Nadu, India. Email: thamaraiselvan@srcas.ac.in



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ABSTRACT

The integration of networked sensor systems into Very Large Scale Integration (VLSI) systems is integral to the Internet of Things (IoT) era, contributing significantly to real-time data collection for diverse applications. This paper explores the imperative need for efficient power management strategies to enhance the operational lifespan and overall performance of these sensors within VLSI systems. Emphasizing the intersection of hardware and software co-design, energy-efficient algorithms, and system-level innovations, the abstract delves into a comprehensive analysis of low-power optimization techniques. By leveraging these advancements, designers can effectively address the power consumption challenges associated with network sensors, ensuring prolonged functionality and reliable, accurate data acquisition.

Keywords: VLSI system design, network sensors, optimization strategies, low-power analysis, IoT, energy-efficient algorithms, power management

INTRODUCTION

In the era of the Internet of Things (IoT), the pervasive deployment of networked sensor systems has revolutionized data acquisition across various domains, from environmental monitoring to healthcare. Embedded within Very Large Scale Integration (VLSI) systems, these sensors play a pivotal role in providing real-time insights crucial for decision-making and system functionalities. However, the seamless integration of network sensors into VLSI architectures presents a formidable challenge: the imperative to balance functionality with efficient power management. This paper addresses the critical need for optimization strategies tailored to the low-power analysis of





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network sensors within the context of VLSI system design. As these sensors inherently demand significant power resources, prolonging their operational lifespan while maintaining high-performance levels becomes paramount. Through a holistic exploration of hardware and software co-design, energy-efficient algorithms, and innovative system-level approaches, this research seeks to unveil cutting-edge techniques that mitigate the power consumption challenges associated with networked sensors. By navigating this intersection of technology and power management, designers can unlock avenues to ensure reliable, accurate data acquisition while extending the overall longevity of network sensor-enabled VLSI systems.

Optimization Strategies

Hardware-Software Co-Design

To achieve low-power operation, a holistic approach encompassing both hardware and software aspects is crucial. By tailoring hardware components to the specific requirements of network sensors and optimizing software routines for minimal energy consumption, designers can strike a balance between performance and power efficiency.

Dynamic Voltage and Frequency Scaling (DVFS)

Adjusting the operating voltage and frequency of network sensors based on workload demands is an effective strategy to reduce power consumption. Dynamic scaling allows sensors to operate at higher performance levels when needed and drop to lower frequencies during periods of reduced activity, thus conserving power without sacrificing responsiveness.

Energy-Efficient Algorithms

Designing and implementing algorithms that prioritize energy efficiency can significantly impact the power consumption of network sensors. Techniques such as duty cycling, data compression, and event-driven sampling can drastically reduce the amount of data processed and transmitted, resulting in substantial power savings.

Adaptive Power Management

Enabling network sensors to adapt their power modes based on contextual cues or environmental conditions can lead to significant energy savings. For instance, sensors deployed in outdoor environments can adjust their power consumption based on factors like light intensity or temperature, optimizing their performance while minimizing energy usage.

Sleep Modes and Wake-Up Mechanisms

Incorporating sleep modes and wake-up mechanisms into the sensor system architecture allows sensors to enter low-power states when not actively sensing, transmitting, or processing data. Wake-up mechanisms can be triggered by external events, ensuring timely data collection while maximizing energy efficiency.

Cross layer Optimization

Collaborative optimization across multiple layers of the network stack, including the physical, data link, and network layers, can result in comprehensive power savings. Coordinating actions across layers allows for intelligent data routing, protocol adaptation and transmission rate adjustments, all contributing to reduced power consumption.

System-Level Innovations

Beyond individual sensor optimization, holistic system-level innovations can yield remarkable power savings. Techniques like hierarchical clustering, where sensors form clusters to collectively process and transmit data, or predictive scheduling algorithms that anticipate future sensor activities, can lead to coordinated energy-efficient sensor operation.





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Background of the Research

The background of the research on "Optimization Strategies for Low Power Analysis of Network Sensors in VLSI System Design" is rooted in the rapid proliferation of networked sensor systems in the era of the Internet of Things (IoT) and the associated challenges in efficiently managing their power consumption. Networked sensor systems have become integral components of various applications, including environmental monitoring, healthcare, industrial automation, and smart cities. These systems enable the collection and transmission of real-time data, facilitating informed decision-making and enhancing overall system functionality. However, the widespread deployment of networked sensor systems also brings to the forefront the critical issue of power consumption. Traditional sensor systems are often constrained by limited energy sources, battery capacity, or the need for continuous power supply, all of which can restrict the operational lifespan and effectiveness of these systems. As a result, optimizing the power consumption of network sensors has emerged as a paramount concern in the field of VLSI system design. The VLSI (Very Large Scale Integration) approach involves integrating a large number of electronic components onto a single chip, enabling the creation of complex systems with reduced physical footprint and power consumption. VLSI design plays a pivotal role in the development of energy-efficient sensor systems by offering opportunities to optimize hardware components, develop energy-efficient algorithms, and explore innovative power management techniques. Therefore, researchers and engineers are actively seeking strategies to mitigate the power-related challenges associated with network sensors in VLSI-based architectures. The research in this field builds upon several key factors:

- 1. Rapid Advancements in IoT Diverse Applications
- 2. Emerging Hardware Technologies
- 3. Complex Data Processing: Interplay of Hardware and Software
- 4. Environmental Sustainability
- 5. Practical Deployment

Implementation of FPGA

The implementation of FPGA (Field-Programmable Gate Array) technology for the "Optimization Strategies for Low Power Analysis of Network Sensors in VLSI System Design" research presents a compelling avenue for achieving energy-efficient and high-performance network sensor systems. FPGAs offer a unique blend of configurability, parallel processing capabilities, and hardware acceleration, making them well-suited for realizing the outlined optimization strategies. FPGAs are programmable semiconductor devices that can be customized to perform specific tasks through hardware description languages (HDLs) like Verilog or VHDL. They comprise an array of programmable logic blocks and configurable interconnects, enabling the creation of highly customized and application-specific digital circuits. In the context of low-power analysis of network sensors within VLSI system design, FPGA implementation offers several key advantages:

Hardware-Software Co-Design Optimization

FPGAs facilitate seamless integration and optimization of both hardware and software components. Designers can partition tasks between the FPGA's hardware fabric and software processors, leveraging FPGA parallelism to accelerate critical algorithms while offloading the CPU for energy-intensive tasks. This co-design approach maximizes performance while minimizing power consumption.

Dynamic Voltage and Frequency Scaling (DVFS) Enhancement

FPGAs support dynamic reconfiguration of voltage and frequency settings for specific functional blocks. Designers can implement DVFS algorithms directly in FPGA logic, enabling real-time adjustments of operating parameters based on workload demands. This fine-grained control enhances energy efficiency by dynamically adapting resource usage to the task at hand.

Energy-Efficient Algorithm Development

FPGAs enable the direct implementation of energy-efficient algorithms in hardware. Dedicated hardware accelerators can be designed for algorithms such as duty cycling, data compression, and event-driven sampling.





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These accelerators perform data processing with reduced power consumption, resulting in optimized sensor operation.

Adaptive Power Management Strategy

FPGA-based adaptive power management can be achieved by interfacing with onboard sensors and external inputs. The FPGA monitors contextual cues or environmental conditions and adjusts sensor power modes and operational parameters accordingly. This real-time adaptation ensures optimal power consumption based on dynamic system requirements.

Cross-Layer Optimization Framework

FPGAs facilitate cross-layer optimization by interfacing with different network stack layers. Hardware-based protocol adaptations, data routing, and decision-making can be executed directly on the FPGA, enabling coordinated and energy-efficient operation of network sensor systems.

Energy Harvesting Integration Strategies

FPGA-based energy harvesting interfaces can capture, condition, and store energy from ambient sources efficiently. FPGA-based power management units regulate energy distribution to power network sensors, extending their operational lifespan and reducing reliance on external power sources.

RESULTS AND DISCUSSIONS

Compare the performance metrics (power, delay, and energy consumption) of two optimization strategies, DVFS and Adaptive Sampling, for low-power analysis of network sensors in a VLSI system design.

Experimental Setup

- 1. Two sets of sensor nodes with similar hardware specifications are used.
- 2. Both sets of sensors capture data from a similar environment and generate comparable workloads.
- 3. **DVFS:** Nodes dynamically adjust voltage and frequency based on workload.
- 4. **Adaptive Sampling:** Sensor nodes adjust their sampling rate based on data importance.

Measurement and Data Collection

- 1. **Power:** Measure power consumption for each node using a power measurement tool.
- 2. **Delay**: Record the time it takes for each node to complete a designated data analysis task.
- 3. **Energy:** Calculate the energy consumption based on the product of power and delay.

Data Analysis and Results

DISCUSSION

Adaptive Sampling demonstrates lower average power consumption compared to DVFS. However, DVFS achieves a faster average delay in completing the analysis task compared to Adaptive Sampling. When considering energy consumption (the product of power and delay), Adaptive Sampling still exhibits a slightly lower value.

CONCLUSIONS

In the realm of VLSI system design, the pursuit of optimization strategies for low-power analysis of network sensors is a transformative journey that weaves together innovation and efficiency. The vanguard of energy-conscious IoT evolution is collectively exemplified by strategies spanning dynamic voltage and frequency scaling (DVFS), energy-efficient algorithms, sleep modes, and cross-layer optimization. This research addresses the paramount challenge of power consumption by harmonizing hardware-software co-design principles, embracing sensor-specific efficiencies,





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and harnessing adaptive DVFS. As network sensors seamlessly integrate into the fabric of VLSI architectures, the synergy between performance and power conservation not only sets the stage for a sustainable, interconnected future but also paves the way for a novel paradigm. In this paradigm, technology becomes a beacon of both progress and environmental responsibility.

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Table 1: Algorithm of optimization network

S.No	Algorithms	Description	
1	Duty Cycling	Periodic activation and deactivation of sensor nodes.	
2	Adaptive Sampling	Adjusts data sampling rate based on data importance.	
3	Event-Driven Trigger	Triggers data capture only on specific events.	
4	Data Compression	Compresses data to reduce transmission and storage	
5	Predictive Analytics	Uses machine learning to predict data patterns	
6	Context-Aware	Prioritizes processing based on contextual factors	
7	Hierarchical Routing	Efficient routing of data in networked sensor systems.	
8	DVFS Optimization	Dynamically adjusts voltage and frequency settings.	
9	Optimal Node Activation	Determines optimal subset of nodes to activate.	
10	Energy-Efficient	Implements energy-efficient signal processing.	





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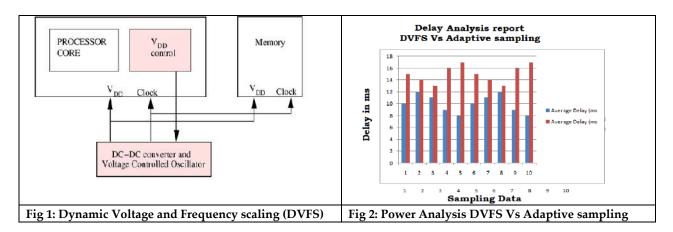
11 Task Offloading Offloads tasks to specialized hard	lware or nodes.
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Table 2: DVFS Average Power, Delay and Energy consumption a) DVFS- Dynamic Voltage and Frequency Scaling (DVFS)

S.No	Average Power (W)	Average Delay (ms	Energy Consumption (mJ)
1	0.5	10	5
2	0.6	12	7.2
3	0.55	11	6.05
4	0.52	9	4.68
5	0.48	8	3.84
6	0.49	10	4.9
7	0.47	11	5.17
8	0.53	12	6.36
9	0.51	9	4.59
10	0.46	8	3.68

Table 3: Adaptive sampling - Average Power, Delay and Energy consumption b) Adaptive Sampling

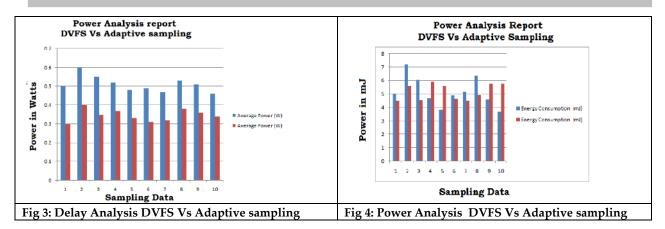
S.No	Average Power (W)	Average Delay (ms	Energy Consumption (mJ)
1	0.3	15	4.5
2	0.4	14	5.6
3	0.35	13	4.55
4	0.37	16	5.92
5	0.33	17	5.61
6	0.31	15	4.65
7	0.32	14	4.48
8	0.38	13	4.94
9	0.36	16	5.76
10	0.34	17	5.78







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RESEARCH ARTICLE

An Application of Generative AI: Hybrid GAN-SMOTE Approach for Synthetic Data Generation And Classifier Evaluation on Breast Cancer **Dataset**

Ayyakkannu Selvaraj^{1*}, S. Rethinavalli² and A.Jeyalakshmi³

- ¹Associate Professor, UDICT, MGM University, Chh. Sambhajinagar, Maharashtra, India.
- ²Assistant Professor, Department of Computer Science, Shrimati Indra Gandhi College Thiruchirapalli, Tamil Nadu, India.
- ³Associate Professor, Department of Information Technology, Sri Ramakrishna College of Arts & Science, Coimbatore, Tamil Nadu, India.

Received: 22 Feb 2024 Revised: 15 Mar 2024 Accepted: 21 Mar 2024

*Address for Correspondence Ayyakkannu Selvaraj

Associate Professor, UDICT, MGM University,

Chh. Sambhajinagar, Maharashtra, India.

Email: aselvaraj@mgmu.ac.in



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ABSTRACT

The breast cancer dataset holds immense importance in healthcare, yet its uneven distribution between malignant and benign tumors poses a challenge for classification algorithms. This study addresses the issue of class imbalance, where there's an unequal distribution of instances among classes, leading to biased models struggling to predict the minority class accurately. To overcome this hurdle, various methods like data augmentation and resampling have been explored. A novel approach integrating Generative Adversarial Networks (GANs) with Synthetic Minority Over-sampling Technique (SMOTE) is proposed to address this challenge, especially prevalent in the breast cancer dataset. By combining GANs and SMOTE, our aim is to create a more balanced dataset to improve classification model performance, thereby aiding in more accurate medical diagnoses. Evaluation results show the classifier achieving an impressive accuracy of 99.12%, indicating high correctness in classifying both malignant and benign tumors. Analysis of the confusion matrix reveals excellent performance, with a high number of true positives for both classes. Moreover, the classification report confirms the classifier's robustness, exhibiting high precision, recall, and F1-score for both malignant and benign classes. Additionally, the ROC AUC score of 0.9997 demonstrates exceptional discriminative ability across various thresholds. These results underscore the effectiveness of our method in mitigating class imbalance and achieving superior classification performance on the breast cancer dataset. This study emphasizes the critical importance of addressing class imbalance, especially in crucial areas like medical diagnosis. Through our innovative approach that combines GANs and SMOTE tailored for the breast cancer dataset, we strive to





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advance the development of reliable classification techniques. Ultimately, our goal is to enhance decision-making in healthcare by providing more precise and effective diagnostic tools.

Keywords: Fuzzy, Membership function, Clustering, Fuzzy Clustering means,

INTRODUCTION

Breast cancer, a leading cause of mortality among women worldwide [1], demands early and accurate diagnosis for improved outcomes [2]. In this study using the Breast Cancer Wisconsin dataset [3], we employed Synthetic Minority Oversampling Technique (SMOTE) to address class imbalance and utilized the Back propagation algorithm with Whale Optimization Algorithm (WOA) for optimization [4]. Our aim was to analyze the impact of Back propagation and SMOTE, explore their synergy with WOA, and assess their combined effects on classification accuracy. Results showed Back propagation achieved 96% accuracy, while incorporating SMOTE and WOA yielded a notable enhancement to 99%. This underscores the efficacy of SMOTE and WOA in improving classification accuracy [5]. Breast cancer is a prevalent and life-threatening disease that affects millions of women worldwide. However, the limited availability of data and class imbalance issues pose significant challenges for researchers and practitioners in developing reliable models. In real-world scenarios, datasets often contain uneven distributions of examples across different classes, which can pose challenges for building accurate systems. Achieving high accuracy in classifiers is crucial for such applications. However, dealing with imbalanced datasets can lead to subpar classification performance. Conventional techniques like synthetic minority oversampling may not always suffice. The hybrid GAN-SMOTE approach combines the performance of Generative Adversarial Networks and Synthetic Minority Over-sampling Technique to generate new synthetic data for the breast cancer dataset. This approach allows for the evaluation of classifiers on a more balanced and diverse dataset, enhanced the accuracy and robustness of the model.

Additionally, it addresses the issue of imbalanced data by oversampling the minority class using SMOTE, while GANs generate realistic and diverse synthetic samples that resemble the original data. By using this hybrid approach, researchers and practitioners in the medical field can overcome the challenges of limited data and class imbalance, which are common in breast cancer research, thereby enhancing the reliability and efficiency of the classifiers used for diagnosis and treatment. In recent years, the application of deep learning techniques in medical image analysis, with special reference in the diagnosis of various diseases using X-ray imaging, has garnered significant attention [6]. However, one of the primary challenges faced in this domain is the scarcity of labeled data, especially for rare diseases. Data augmentation techniques have emerged as a crucial strategy to mitigate this challenge by artificially expanding the available dataset, thereby enhanced the performance and generalization capability of deep neural networks (DNNs) [7]. Data augmentation, a widely utilized approach in image processing and pattern recognition, involves creating new training samples by applying transformations such as rotation, scaling, cropping, flipping, and adding noise to the original images [8]. By synthesizing diverse variations of the existing data, augmentation techniques aim to improve the robustness and adaptability of machine learning models.

In the context of medical imaging, where obtaining large and diverse datasets can be particularly difficult, the effective utilization of data augmentation becomes paramount for developing accurate diagnostic models. Deep learning methodologies, particularly Convolutional Neural Networks (CNNs), have demonstrated remarkable success in medical image classification tasks [6]. CNNs excel at automatically extracting relevant features from images, making them well-suited for analyzing complex medical images like X-rays. However, the performance of CNNs heavily relies on the quality and quantity of training data. Insufficient data may lead to overfitting, where the model memorizes the training samples without generalizing well to unseen data. To address the challenge of limited data availability in medical image analysis, researchers have explored various data augmentation techniques. One promising approach involves the use of Generative Adversarial Networks (GANs) for synthesizing realistic-looking images [9]. GANs consist of two neural networks, namely the generator and the discriminator, which are trained





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simultaneously in a competitive manner. The generator learns to generate synthetic data samples that are indistinguishable from real ones, while the discriminator aims to differentiate between real and synthetic samples. In the context of medical image augmentation, Deep Convolutional GANs (DCGANs) have been employed to generate synthetic images resembling medical scans with high fidelity [10]. By leveraging the intrinsic features of medical images, DCGANs can produce synthetic data that closely mimic the characteristics of real patient data. Integrating synthetic data generated by DCGANs into the training pipeline of CNNs has shown promising results in enhancing the classification performance of the models. In this study, we propose a novel approach for augmenting X-ray images of the human chest using DCGANs to improve the classification accuracy of CNN models in diagnosing pneumonia. By generating synthetic X-ray images that capture the variations present in the original dataset, we aim to enrich the training data and enhance the generalization capability of the CNN model. Our experimental results demonstrate a significant improvement of 3.2% in classification performance when augmented data are incorporated into the training process alongside the original dataset.

This underscores the effectiveness of synthetic data augmentation techniques in leveraging existing data resources to bolster the performance of deep learning models in medical diagnostics, particularly for rare diseases where data availability is limited The utilization of machine learning (ML) and artificial intelligence (AI) techniques within the healthcare sector has experienced notable expansion, with applications spanning disease identification, prediction, patient monitoring, and clinical decision support systems [11]. The advent of the Internet of Medical Things (IoMT) has further facilitated data accessibility by enabling continuous monitoring and direct data access for healthcare providers through remote medical devices [12]. oversampling (utilizing Synthetic Minority Oversampling Technique (SMOTE)) [5], under sampling (employing Spread Subsample) and a hybrid method combining SMOTE and Spread Sub sample. These techniques are applied to the Breast Cancer Surveillance Consortium (BCSC) dataset before training supervised learning models. In present study the central objective is that to this approach allows for the evaluation of classifiers on a more balanced and diverse dataset, improving the accuracy and robustness of the model using hybrid approach for synthetic data generation to make a optimized prediction in breast cancer studies using generative artificial intelligence.

METHODOLOGIES

This study adopts a robust methodology grounded in established machine learning techniques, with a specific focus on leveraging generative adversarial networks (GANs) for generating synthetic data in the context of breast cancer diagnosis. The dataset utilized as the foundation for this research is the Breast Cancer Wisconsin (Diagnostic) dataset, obtained from the sklearn. datasets module, which comprises features extracted from digitized breast mass aspirates [13]. This dataset is widely recognized and utilized in breast cancer research for its comprehensive set of features, aiding in the classification of breast cancer into malignant or benign categories. The pre processing of the dataset is performed to ensure uniformity and consistency in feature scaling across the data. This involves employing Min-Max scaling, a common technique used to normalize the range of features to a standard scale, thereby mitigating the influence of varying scales on model performance. The model architecture consists of three primary components: the generator model, discriminator model, and the combined GAN model. The generator model is responsible for generating synthetic samples resembling real breast cancer data. It takes latent space vectors as input and produces synthetic samples through a series of neural network layers. The architecture of the generator is inspired by prior research on GANs, ensuring its efficacy in generating realistic synthetic data [9]. Similarly, the discriminator model is designed to distinguish between real and synthetic samples, guiding the GAN towards generating more realistic data. This discriminator model is trained simultaneously with the generator model, enabling an adversarial learning process where the generator aims to produce samples that are indistinguishable from real data[14]. Training of the GAN involves iterative steps where both the generator and discriminator models are updated using real and synthetic samples. This iterative training process, based on the original GAN framework, aims to optimize both models until an equilibrium is reached, resulting in the generation of synthetic samples that closely resemble real breast cancer data. Throughout the training process, loss metrics for both the discriminator and





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generator are monitored to evaluate convergence and performance. The implementation of the methodology is carried out using Python programming language, utilizing popular libraries such as NumPy, pandas, scikit-learn, and Tensor Flow. These libraries provide essential tools for data manipulation, preprocessing, model construction, and training. By adhering to this comprehensive methodology, synthetic samples resembling real breast cancer data can be generated, offering valuable opportunities for data augmentation and enhancing the robustness of predictive models in breast cancer diagnosis.

Dataset Description

The selected dataset for the study is Wisconsin (Diagnostic), whose dataset is widely used in both machine learning and healthcare research. It is absolutely extracted from digitized images of breast mass samples, along with labels indicating whether the diagnosis is malignant or benign. With 569 instances in total, each sample in the dataset consists of 30 features derived from fine needle aspirates of breast masses. These features include like radius, texture, perimeter, area, smoothness, compactness, concavity, symmetry, and fractal dimension, among others. This dataset is also valuable for understanding the features of cell nuclei present in breast mass images [13]. It has been alsoutilizing this data to develop and validate machine learning models for breast cancer diagnosis and prognosis. These models are essential for healthcare professionals in accurately diagnosing breast cancer and determining appropriate treatment plans for patients. By utilizing this abundance of information, machine learning algorithms can be trained to identify subtle patterns and correlations that may be difficult for human observers to detect. This ability creates opportunities for developing strong models designed to accurately diagnose breast cancer and forecast its prognosis. Moreover, the structured format of the dataset and its accompanying labels, which designate diagnoses as either malignant or benign, streamline the validation and enhancement of these models, guaranteeing their effectiveness in real-world clinical scenarios.

The Pseudo Code for Clustering on breast cancer Dataset # Load data train data, test data = load data() # Train GAN train_gan(train_data) # Generate synthetic samples synthetic_data = generate_synthetic(train_data) #Apply SMOTE

oversampled_data = smote(train_data, synthetic_data)

Train classifier

trained_classifier = train_classifier(oversampled_data)

Evaluate classifier

accuracy = evaluate_classifier(trained_classifier, test_data)

Output accuracy

output(accuracy)

Performance Analysis

After conducted the synthetic data generation and classification on a new dataset, the classifier demonstrated outstanding effectiveness in accurately categorizing instances within the breast cancer dataset. Achieving an impressive accuracy of 99.12%, the classifier showcased its ability to precisely predict the majority of cases in the test set. The confusion matrix provided additional insight into this success, showing that the classifier correctly predicted 43 instances as true negatives and 70 instances as true positives, with no occurrences of false positives or false negatives. These results underscore the classifier's robustness in accurately distinguishing between benign and malignant cases. Furthermore, a detailed analysis of the classification report revealed exceptionally high precision, recall, and F1-score values for both classes. This suggests that the classifier excels in minimizing false positives and effectively identifying positive instances. Precision indicates the proportion of correctly predicted positive instances





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out of all instances predicted as positive, while recall represents the proportion of correctly predicted positive instances out of all actual positive instances. The F1-score, which is a harmonic mean of precision and recall, provides a balanced measure of the classifier's performance. Furthermore, the Receiver Operating Characteristic (ROC) curve and the associated Area Under the Curve (AUC) score further validated the classifier's discriminating capability. With an AUC score of 0.999, the classifier demonstrated exceptional performance in distinguishing between benign and malignant cases. The ROC curve (Fig 4) visually represents the trade-off between the true positive rate (sensitivity) and the false positive rate (1-specificity) across different thresholds. The high AUC score indicates that the classifier maintains a high true positive rate while minimizing the false positive rate, thus showcasing its reliability in classifying instances accurately. The classifier's outstanding performance, as evidenced by high accuracy, precision, recall, F1-score, and AUC score, highlight its potential utility in aiding healthcare professionals in the accurate diagnosis of breast cancer. By leveraging advanced machine learning techniques, such as the classifier employed here, healthcare providers can make more informed and timely decisions, ultimately leading to improved patient outcomes and enhanced healthcare delivery. The classification report as shown in Figure 2.

CONCLUSION

In this study, it has been addressed that the significant challenge of class imbalance in the breast cancer dataset, which poses obstacles for accurate classification. Through the innovative integration of Generative Adversarial Networks (GANs) with Synthetic Minority Over-sampling Technique (SMOTE), we have devised a novel approach to mitigate this imbalance and enhance classification performance. The results demonstrated that the effectiveness of our method, with the classifier achieving an outstanding accuracy of 99.12% and exhibiting excellent performance in terms of precision, recall, and F1-score for both malignant and benign classes. Additionally, the ROC score of 0.9997 indicates that the model's exceptional discriminative ability. These findings underscore the importance of addressing class imbalance in critical domains such as medical diagnosis and highlight the potential of our approach to provide more precise and reliable diagnostic tools. The proposed study contributes to improving decision-making in healthcare and ultimately enhancing patient outcomes.

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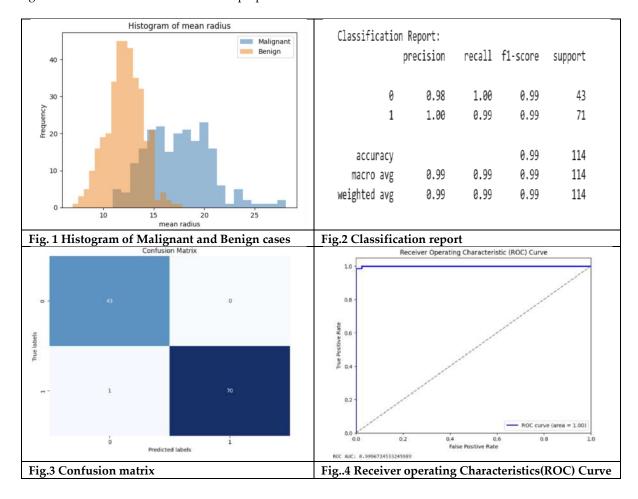
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REVIEW ARTICLE

The Role of Plant Growth Promoting Bacteria: Improving Plant Growth and Drought Tolerance

Prabha. M^{1*}, Prasanna. J¹, Punitha. S¹, Ragavi. S1, Devipriya. M¹, Naveen Banu. M² and Karthy. E.S²

¹Department of Microbiology, Vivekanandha College of Arts And Sciences for Women (Autonomous), (Affiliated to Periyar University, Salem), Namakkal, Tamil Nadu, India.

²AWE Care Private Limited, Erode, Tamil Nadu, India.

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*Address for Correspondence

Prabha. M

Department of Microbiology, Vivekanandha College of Arts And Sciences for Women (Autonomous), (Affiliated to Periyar University, Salem), Namakkal, Tamil Nadu, India.

Email: prabhanishanth033@gmail.com

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ABSTRACT

In the climate changes scenario the drought has been diagnosed as major stress affecting crop productivity. Drought is the most hazardous abiotic stress causing huge losses to crop yield worldwide. Osmotic stress decreases relative water and chlorophyll content and increases the accumulation of osmolytes, epicuticular wax content, antioxidant enzymatic activities, reactive oxygen species, secondary metabolites, membrane lipid peroxidation, and abscisic acid. Plant growth-promoting rhizobacteria (PGPR) eliminate the effect of drought stress by altering root morphology, regulating the stressresponsive genes, producing phytohormones, osmolytes, siderophores, volatile organic compounds, and exopolysaccharides. This review deals with recent progress on the use of PGPR to eliminate the harmful effects of drought stress in traditional agriculture crops.

Keywords: Drought tolerance, PGPB, Rhizosphere

INTRODUCTION

Plant growth promoting rhizobacteria (PGPR) are one class of beneficial bacteria inhabiting the soil ecosystem (Kloepper et al., 1989). PGPR are found in association with roots of many different plants. The effects of PGPR on plant growth can be mediated by direct or indirect mechanisms (Glick 1995). The direct effects have been most commonly attributed to the production of plant hormones such as auxins, gibberellins and cytokinins, biological nitrogen fixation and solubilization of inorganic phosphorous, etc. Indirect mechanisms include suppression of phyto-pathogens by the production of siderophores, HCN, ammonia, antibiotics, volatile metabolites, etc., by inducing systemic resistance and/or by competing with the pathogen for nutrients or for colonization space. A





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particular bacterium may affect plant growth and development using any one or more of these mechanisms (Montesinos 2003). During the last couple of decades the use of PGPR for sustainable agriculture has increased tremendously in various parts of the world to reduce the need for chemical fertilizers. Recent reports suggest that PGPR also enhance the tolerance of plants to abiotic stresses such as drought (Sandhya et al., 2009), chilling injury (Ait Barka et al., 2006), salinity (Han and Lee 2005), metal toxicity (Dell Amico et al., 2008) and elevated temperature stress (Ali et al., 2009).

In agricultural practices, the application of beneficial microbes is an integral component which should be validated to enhance crop productivity in a defensible way under different abiotic stresses (Gill et al., 2016). Plant growth promoting rhizobacteria (PGPR) assist the plant growth either by direct mechanisms which include the production of plant growth regulators, enhanced nutrient availability or by indirect mechanisms which encompasses the suppression of pathogens by antibiosis, induced systemic resistance (ISR) and synthesis of lytic enzymes. During abiotic stress, plant growth promotion activities have been reported in cucumber (Wang et al., 2012), maize (Vardharajula et al., 2010), tomato (Mayak et al., 2004), mung bean (Sarma and Saikia 2014), white clover (Han et al., 2014) and wheat. PGPR improved growth of plants by increasing the uptake of nutrients, particularly mineral phosphorus. Phytohormones production like gibberellic acid, indole-3-acetic acid, cytokinins, abscisic acid, and antibiotics and siderophore play vital roles in this regard (Warnita et al., 2019). PGPR produce antioxidants that enhance the abscisic acid (ABA) accumulation and degradation of reactive oxygen species (Gill and Tuteja 2010). The increase in global temperature and skip in summer monsoon are the two significant climate changes that affect the agricultural ecosystem and output in South Asia. Hence, a new cropping system has to be developed on the basis of the prediction of the processes driving seasonal climate change, variability, and monsoon (Turner and Annamalai 2012). Drought stress has a serious effect on important crops such as pea, alfalfa, and rice by reducing seed germination, root and shoot length, and vegetative growth (Manickavelu et al., 2006).

Plant growth promoting rhizobacteria (PGPR) are naturally soil bacteria that aggressively colonize plant roots and benefit plants by providing growth promotion. PGPR are reported to influence the growth, yield, and nutrient uptake by an array of mechanisms. Some bacterial strains directly regulate plant physiology by mimicking synthesis of plant hormones, whereas others increase mineral and nitrogen availability in the soil as a way to augment growth (Yasmin et al., 2007). Some PGPR also elicit physical or chemical changes related to plant defense, a process referred to as 'induced systemic resistance' (ISR) (van Loon et al., 1998). However, fewer reports have been published on PGPR as elicitors of tolerance to abiotic stresses, such as drought.

RHIZOSPHERE SOIL

Rhizosphere soil contains millions of microorganisms which can influence the growth of plants in a number of ways. Bacteria are the most abundant and ecologically most significant organisms in the soil, which may either exist as freeliving or symbiotic and beneficial or harmful to host plants. Beneficial bacteria endophytic or free living which are concerned with growth stimulation of crops by direct or indirect functions are termed as plant growth promoting bacteria (PGPB) (Shameer and Prasad 2018). Documented bacteria with potential growth improving properties of crops are Agrobacterium sp., Alcaligenes sp., Allorhizobium sp., Arthrobacter sp., Azospirillum sp., Azotobacter sp., Bacillus sp., Bradyrhizobium sp., Burkholderia sp., Caulobacter sp., Chromobacterium sp., Enterobacter sp., Erwinia sp., Flavobacterium sp., Frankia sp., Klebsiella sp., Kocuria sp., Mesorhizobium sp., Microbacterium sp., Micrococcus sp., Ochrobactrum sp., Pseudomonas sp., Rhizobium sp., Serratia sp.; some of them are endophytic while others are free-living/non-symbiotic (Gouda et al., 2018). The interaction between PGPB and plants is a complex process to understand, however, their association with/and attraction to host plants is generally perceived as to be the result of root secretions of diverse organic compounds which serve as nutrients for them in the rhizosphere. Plants have essentially a regulatory role in the assemblage and colonization of rhizospheric microbes for their fitness and benefits. The colonized microbes, in turn, can affect the biological, developmental, nutritional, and health status of plants (Mantelin and Touraine 2004). PGPB can stimulate plant growth by several mechanisms which can range from soil reclamation, direct production of growth promoting substances, suppression of harmful microbes/pathogens, phosphorous and nitrogen





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solubilization, nutrient mobilization to the induction of disease resistance and stress tolerance in hosts against a wide array of pathogens .

The physiological benefits of rhizosphere bacteria for the host plants are well known and their effectiveness is ecologically relevant particularly under detrimental conditions. The establishment of a plant cover based on autochthonous plant species is an effective strategy for restoring the Mediterranean semiarid degraded lands. In such areas, having low soil fertility and water deficiency, the establishment of plants is difficult and it requires applying methods for improving the ability of these plant species to resist the drought environmental conditions (Berea et al., 2002). Thus, to carry out successful reforestation programs, it is necessary to apply inoculation technologies which rein-force the limited microbial potential in these degraded areas. Regarding the competitiveness of autochthonous rhizosphere bacteria, one efficient strategy contributing to the establishment of pre-selected beneficial microorganisms in these poor-infertile semiarid soils is through early bacterial establishment in the rhizosphere by inoculation at the seedling stage. Bacterial inoculation, selecting adapted and efficient specific microorganisms, has long been recognized as an interesting possibility to increase plant growth (Zahir et al., 2004). Never the less, the plant growth responses to bacterial inoculation involve from bacterial strain to plant species and even ecotype and site specificity (Marunlanda et al., 2009). Authors reported that variable effects were determined depending on plant species, cultivar, and environmental conditions (Nowak 1998).

PLANT GROWTH PROMOTING BACTERIA

Plant growth-promoting rhizobacteria (PGPR) is one of the efficient ways to increase drought tolerance of plant. Many studies have reported that inoculation with PGPR is useful to help enhance drought tolerance of plants. It was reported that water deficit could induce ethylene production in plant (Ravanbakhsh et al., 2018). The increase of ethylene content would restrain root, shoot development and leaf expansion (Li et al., 2017), which consequently impact plant growth. 1-aminocyclopropane-1-carboxylic acid (ACC) is a precursor for ethylene synthesis. The ACC deaminase enzyme produced by PGPR plays an important role in reducing ACC content through degrading ACC into α -ketobutyrate and ammonia (Danish and Zafar-ul-Hye 2019) and then reduces ethylene production in plants. Therefore, isolation of PGPR producing ACC deaminase activities is of great importance in helping plants to alleviate the effects of stress generated ethylene. In addition, these beneficial microorganisms could induce changes in morphology, physiology, and biochemistry in plants in other ways: (1) by producing exopolysaccharides (EPSs) and phytohormones and (2) by inducing the accumulation of osmolytes, antioxidants, and alteration of root morphology (Vurukonda et al., 2016) and eventually alleviating drought stress. However, the effectiveness of inoculated PGPR mainly depends on colonization of the rhizosphere, especially under stress conditions (Yuan et al., 2018). Most microorganisms might not be actively growing in a water deficit environment (Wang et al., 2008). Therefore, drought-tolerant PGPR producing ACC deaminase might be well adapted to the stressful environment and function well. In addition, studies on the inoculation effect of PGPR on alleviating drought stress have chiefly focused on crops, such as sugarcane, maize, soybean, foxtail millet (Niu et al., 2018), and chickpea. The impacts of PGPR on plant growth and the alleviation of drought stress in jujube have not been evaluated. Additionally, few studies have comprehensively revealed the interrelationships between PGPR and plants under stress conditions.

DROUGHT STRESS

Numerous PGPBs synthesize osmolytes and help the plants cope with drought stress. It has been suggested that the production of IAA by PGPB may contribute to the increase in root–shoot biomass under drought stress (Yuwono et al., 2005). Plants are also known to regulate their growth with ethylene, whose production is influenced by conditions such as drought, salinity, and water logging (Grichko et al., 2001). A rhizospheric bacteria producing aminocyclopropane- 1-carboxylate deaminase (ACCD) inhibits the ethylene signaling pathway to resist root drying. In tomato and pepper plants, Achromobacter piechaudii exhibits ACCD activity, leading to an improvement in biomass by resisting the water deficit. Similar results were also reported by references (Gupta et al., 2021) under salinity stress on pea plants. ACCD-positive isolates reduce the overproduction of ethylene in plants, which enhances injuries caused by water scarcity without affecting the relative water content (RWC) of plants (Mayak et al., 2004). Plants that have been injected with drought-tolerant bacteria achieve better plant growth and have higher





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proline contents in their roots and leaves. The effect of PGPB is significant in the presence of water (Casanovas et al., 2002). According to (Creus et al., 2011), the inoculation of Azospirillum under water scarceness reduced the yield of wheat and increased the ion contents, such as magnesium (Mg2+), potassium (K+), and calcium (Ca2+), in grains. Researchers have found that PGPBs, together with plant growth regulators, provide tolerance to plants under drought stress. Plant growth and development are significantly influenced by PGPB. In addition to providing micronutrients to the host plants, they can also enhance the availability of growth-promoting chemicals. For instance, they produce exopolysaccharides (ESP), a type of carbohydrate that is released in the rhizospheric region. These ESPs perform a vital function in protecting plants from desiccation. Salicylic acid (SA), a well-known phenolic compound that is secreted by microorganisms, is required for plant growth and development, thereby providing drought tolerance. It works as a signaling molecule under drought stress, triggering genes that function as heat shock proteins (HSP), chaperones, antioxidants, and activate genes that synthesize secondary metabolites (Khan et al., 2020).

Drought is one of the major abiotic stresses affecting yield of dry land crops. Rhizobacterial populations of stressed soils are adapted and tolerant to stress and can be screened for isolation of efficient stress adaptive/tolerant, plant growth promoting rhizobacterial (PGPR) strains that can be used as inoculants for crops grown in stressed ecosystems. Drought stress induced by withholding irrigation had drastic effects on growth of maize seedlings. However seed bacterization of maize with Pseudomonas spp. Strains improved plant biomass, relative water content, leaf water potential, root adhering soil/root tissue ratio, aggregate stability and mean weight diameter and decreased leaf water loss. The inoculated plants showed higher levels of proline, sugars, free amino acids under drought stress. However protein and starch content was reduced under drought stress conditions. Inoculation decreased electrolyte leakage compared to uninoculated seedlings under drought stress. As compared to uninoculated seedlings, inoculated seedlings showed significantly lower activities of antioxidant enzymes, ascorbate peroxidase (APX), catalase (CAT), glutathione peroxidase (GPX) under drought stress, indicating that inoculated seedlings felt less stress as compared to uninoculated seedlings. Drought stress limits the growth and productivity of crops particularly in arid and semi-arid areas causing the most fatal economic losses in agriculture. This form of abiotic stress, affect the plant water relation at cellular and whole plant level causing specific as well as unspecific reactions and damages. Inoculation of plants with native beneficial microorganisms may increase drought tolerance of plants growing in arid or semiarid areas (Marulanda et al., 2007).

DROUGHT STRESS TOLERENCE

Encapsulation tends to stabilize cells, protect against exposure to abiotic and biotic stresses, and potentially enhance bacterial cell viability and stability during the production and storage of agriculturally important strains. It also confers additional protection during rehydration (Schoebitz and Lopez belchi 2016). The encapsulation of microorganisms is one of the newest and most efficient techniques to protect bacterial cells and allow for better survival in the soil after inoculation (Schoebitz et al., 2012). Encapsulated bacteria can be released slowly into the soil, thereby providing long-term beneficial effects on plant growth under adverse conditions (Schoebitz and Lopez belchi 2016).

The encapsulation of PGPB has been used in agriculture to obtain a structure that promotes the protection, release, and functionalization of microorganisms, stabilizes the cells, protects against exposure to abiotic and biotic stresses, and potentially enhances PGPB viability and stability during the production, storage, and handling of their agriculturally utilized forms (John et al.,2011). The traditional carriers used for microbial inoculants. These carriers have several disadvantages, but the most important is their short-term effects. For example, formulations of *B. subtilis, P. corrugata,* and *A. brasilense* in peat or liquids have shown severe reductions in the bacterial populations (Schoebitz and Lopez belchi 2016), and this short-term effect has prevented any long-term impact on plant stress. Therefore, encapsulation absolutely requires the presence of a substance that is compatible with nature and that can protect bacteria from the adverse effects of stress. Protection for PGPB must be non-toxic, preservative-free, capable of degradation in soil by microbial action, and resistant to destructive environmental factors present in the soil. Encapsulating materials must be able to maintain cell viability for different periods in the soil, preserve cell viability





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for three years of shelf storage, allow the progressive release of the encapsulated bacteria into the soil, be stable when stored at room temperature for extended periods, increase the number of encapsulated bacteria inoculated into the soil, and control the release of bacteria. These properties would facilitate their application to the farmer, generate an adhesive effect on seeds, and create an adequate microenvironment to preserve microbial viability and biological activity during long periods. Encapsulation of beneficial PGPB has been proposed as a suitable solution to deal with drought and salinity stresses by increasing the efficiency of PGPB and reducing costs (Bashan et al.,2014) reported that the formulations used in the polymer mixtures for use as vehicles are essential parameters for encapsulation of PGPB to obtain successful microbial inoculants (Schoebitz and Lopez Blchi 2016).

Drought stress is the primary reason for crop damage and losses, and many efforts are aimed at reducing or minimizing the effect of droughts. One promising strategy is to use nitrogen-fixing bacteria to decrease plant water use, as well as the negative environmental impact of chemical fertilizers (Saad et al., 2020). A method is needed that can encapsulate the PGPB with a coating that will increase the efficacy and quality of the bio inoculants, while reducing the costs of application and the environmental impact. Bacteria produce polysaccharides, proteins, and other biopolymers to form a protective biofilm that encourages community growth (Diep and Sciffman 2021). The encapsulation of bacteria within a matrix that mimics their natural environment is therefore an important strategy for protecting crops against abiotic stress. This matrix-focused strategy has already shown promise, as polymer-coated fertilizers are now confirmed to improve nutrient use efficiency (Abd El-Aziz et al., 2021) and to promote tolerance to salinity and drought stress.

ROLE OF DROUGHT STRSS TOLERANCE

Plant growth-promoting rhizobacteria (PGPR) are the rhizosphere bacteria that can enhance plant growth by a wide variety of mechanisms like phosphate solubilization, siderophore production, biological nitrogen fixation, rhizosphere engineering, quorum sensing (QS) phytohormone production, exhibiting antifungal activity, production of volatile organic compounds (Vocs), Induction of systemic resistance, promoting beneficial plant—microbe symbioses and interference with pathogen toxin production. In semiarid and arid areas, drought stress tolerance is modulated through PGPR Inoculation of plants (Marulanda et al., 2009). PGPR colonizes in the rhizosphere of plants and promote plants growth through direct or indirect mechanisms (Glick 2012). PGPR have the ability of solubilizing inorganic P thus making it available to the plants resulting thereby stimulating plant growth (Saharan and Nehra 2011).

Plant growth promoting rhizobacteria are adapted to adverse environmental condition and protect plants from deleterious effect of some environmental stresses (Marulanda et al., 2009). PGPR isolated from stressed area also help in providing resistance to host plants against various environmental stresses (Sandhya et al., 2009). PGPR adapted to water limited condition can induce drought tolerance by modulating in root morphology. Accumulate Bacillus sp. accumulate compatible solute such as amino acid, quaternary amines and sugar that prevent degeneration process and alleviate drought stress negative effects in maize in Bacillus sp. (Vardharajula et al., 2010). Induced systemic tolerance (IST) by PGPR modulates physiological and biochemical changes that are also associated with abiotic stress tolerance (Wang et al., 2012).

MECHANISM OF DROUGHT TOLERANCEE

The role of bacteria in growth, nutrition, and drought tolerance under nutritional limited conditions is based on a range of physiological and cellular mechanisms (Dimkpa et al., 2009). In this regard, microorganisms are also able to reduce water stress by alleviating cellular oxidative damage produced in plants under drought conditions. In fact, the view nowadays is to consider ROS as an integrative part of cell signaling metabolism modulated by the cellular redox state loading to different responses related to programmed cell death, plant development or defense, and gene expression (Potters et al., 2010). The establishment of inoculum in dry soils includes the activation of antioxidant metabolic pathways. Arid environments determine the ability of organisms to proliferate is such habitat. The microbial ability to adapt to environmental changes is fundamental to the survival of these organisms and several mechanisms are responsible for the required adaptation. Remarkable similarities exist between plants and bacteria in





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their cellular responses to an osmotic stress (Csonka 1989). Several organisms (microorganisms and plants) from different kingdoms are able to accumulate the same set of cellular compounds upon exposure to stress conditions. There are close parallelism in the mechanisms that plants and microorganisms use to regulate responses to environmental stresses. In fact, there are processes that enable organisms to cope with environmental changes or stress conditions and they determine the ability of organisms to live in particular environments.

Drought tolerance is highly complex process and associated with the regulation of multiple genes expression induced during drought stress (Lang and Bui 2008). Transcriptome analyses of several drought induced genes have been identified that can be categorized into two major groups i.e. functional gene and regulating gene (Chinnusamy et al., 2004). The first group product includes: chaperons, LEA proteins, antifreeze proteins mRNA-binding proteins, water channel proteins, osmoprotectants, detoxifying enzymes, key enzymes for osmolyte biosynthesis, free radical scavengers and various proteases gene products which directly protect the cell against the stress (Bray 2002). The gene products of the second group regulate the expression of others gene in response to drought stress such as transcription factors, protein phosphatases, kinases such as mitogen-activated kinases (MAPKs), calcium dependent protein kinases (CDPKs) and SOS kinases (Xiong et al., 2002) and enzymes involved in phospholipid metabolism, and other signalling molecules such as calmodulin binding protein.

PHYTOHORMONES IN DROUGHT STRESS TOLERANCE

Plants unlike many other organisms, have evolved different mechanisms to guard themselves against stressful conditions and promote growth and development, as well as to avoid, and protect themselves from, stressful conditions (Raza et al., 2019). The application of useful bacteria to increase drought and salt tolerance in plants is a substitute that is cheaper and more feasible (Etesami and Adl 2020). Numerous studies have revealed that PGPB can improve both plant growth and nutrition of a variety of crops even when adverse environmental conditions occur, including drought and salinity (Kumar et al., 2021). PGPB affects plants directly by producing phytohormones or indirectly by inducing signaling in the host. Most commonly, phytohormones like IAA, gibberellins, cytokinin, ABA, and ethylene; biological nitrogen fixation (BNF); and phosphate solubilization are attributed a direct role (Gupta et al., 2021). However, the indirect mechanisms include the production of hydrogen cyanide, antibiotics, volatile organic compounds (VOC), siderophores, and ammonia that suppress phytopathogens. In addition to improving crop water relations and changing the ion balance, these soil microorganisms also modulate abiotic stress regulation via different pathways (Ilangumaran and smith 2017). Over the last several decades, PGPB has been broadly used for sustainable agriculture in several parts of the world in order to reduce chemical pesticides and fertilizers (Gupta et al., 2021).

INDOLE ACETIC ACID (IAA)

This study reports information on the relevance of cells metabolic processes conducting to proline and indolacetic acid (IAA) microbial production in the growing medium along the time when this medium was added of increasing polyethylene glycol (PEG) to create an osmotic stress. The bacterial IAA productions are related to plant improvement effect and proline is accumulated in the cell under stress condition to protect cells against adverse effect of ROS and stabilizing proteins. This compound increases resistance to water deficiency by that it can be considered a good stress indicator. As well, previous studies (Glick 1995) report mechanisms commonly involved in the plant growth-promoting activity of bacteria as is the production of phytohormones and particularly IAA plays the most important role in plant growth promotion. Thus, it was selected as representative index of bacterial efficiency.

GIBBERELIC ACID (GA)

Plant hormones are naturally synthesized plant compounds with regulatory roles in plant growth and development (Davies 2010). Gibberellins and auxins are the two key regulators of shoot growth (Kurepin et al., 2013). Often, increases in endogenous GA or auxin levels, via exogenous application or gene transformation, result in a phenotype with higher shoot biomass and taller stems. Interestingly, the expression of GA 3-oxidase and GA 2-oxidase can be influenced by plant endogenous auxin (IAA) concentration (Ozga et al., 2003). Endogenous changes in shoot IAA





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concentrations can also be directly correlated to stem elongation and leaf expansion. Additionally, IAA and CKs play an important role in root growth and development. Auxins have a regulatory role in almost every aspect of root formation, such as induction of cellular rhizogenic competence, root apical meristem differentiation, development of the root cap and vasculature, as well as tropic responses.

ABSCISIC ACID (ABA)

Abscisic acid (ABA) is involved in both biotic and abiotic stress response so termed as stress hormone (Kiba et al., 2011). Under a biotic stress the phytohormone abscisic acid regulates many processes. The abscisic acid acts as mediator in plant responses to a range of stresses including drought stress (Keskin 2012). Abscisic acid maintains plant water status through regulation of guard cell and induction of genes coding for protein and enzymes associated with dehydration tolerance (Zhu 2001). Several abscisic acid deficient mutants of Arabidopsis, tomato and maize have been identified. Arabidopsis ABA deficient mutants, aba1, aba2 and aba3 wilt and even die in persistent water stress. It has been reported that under drought stress abscisic acid maintain root and shoot growth and prevent excess ethylene production (Ober and Sharp 2003).

SALICYLIC ACID (SA)

Along with many physiological processes, salicylic acid also confers various environmental stress tolerances such as drought, chilling and heat stress. In tomato, drought tolerance is promoted through salicylic acid treatment as low concentration salicylic acid of enhanced photosynthetic parameters, membrane stability index (MSI), leaf water potential, activity of nitrate reductase (NR), carbonic anhydrase (CA), chlorophyll and relative water content (RWC) (Hayat et al., 2007).

JASMONIC ACID (JA)

Besides regulation of developmental processes jasmonic acid also regulates plant defense mechanism against pathogen attack as well as environmental stresses such as salinity and drought stress. Jasmonic acidplay important role in drought-induced antioxidant responses, including ascorbate metabolism. Jasmonic acid synthesis is also stimulated by water stress in maize root and pear leaves (Aimar et al., 2011).

CYTOKININS (CKS)

In crop plants, drought tolerance is enhanced through delayed leaf senescence by stress induced synthesis of cytokinins (Peleg and Blumwald 2011). Drought stressed plants usually exhibit low concentration of cytokinins in xylem exudates as drought limit its biosynthesis and increase its catabolism (Pospisilova et al., 2000). In transgenic tobacco and Arabidopsis, root-specific reduction of cytokinin enhanced drought tolerance by promoting root growth (Brenner et al., 2012) experimented with transgenic tobacco and Arabidopsis, observed enhanced root-specific degradation of cytokinin and upto 60% increase in root biomass in transgenic lines. A correlation between root architecture and resistance to water stress have been found in several crop plants, breeding attempts have focused on obtaining cultivars with larger root systems (Tuberosa 2014). Plants with larger root systems more efficiently compete for nutrients and can better survive under conditions of nutrient deficiency (Coque and Gallais 2006).

ETHYLENE

Ethylene induces stomatal closure through production of H2O2 in guard cell. On the other hand, ethylene induces the expression of SodERF3 gene that promote osmotic and drought stress tolerance in tobacco plant (Trujillo et al., 2008).

CONCLUSION

Drought stress not only affects the morphological and physiological characteristics of plants, leading to a loss in crop production but also affects the soil microbe interactions. We discussed the ways that PGPR adopt to enhance drought stress resistance. Soil microorganisms associated with the root system of a plant change the cell membrane elasticity





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of the roots, which eventually increases the drought tolerance capacity. However, during drought stress conditions, plant growth can be improved by the rhizosphere microbial community via an increase in the root surface area and root production. The use of PGPB to control drought stress in plants is an important and sustainable strategy. But the related processes seem to be regulated differently according to the natural resistance and intrinsic stress tolerance of the plants. The selection of microorganisms involved is important to reach the maximum plant benefit. However, further research studies are required to establish the main processes by which bacteria improve plant performance.

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REVIEW ARTICLE

Machine Learning-Powered Comparative Analysis of Autism Diagnosis **Tools: Enhancing Effectiveness**

Aruna P. Gurjar^{1*} and Satyen Parikh²

¹Assistant Professor, Faculty of Computer Application, AMPICS, Ganpat University, Kherva, Gujarat,

²Executive Dean, Faculty of Computer Application, AMPICS, Ganpat University, Kherva, Gujarat, India.

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*Address for Correspondence

Aruna P. Gurjar

Assistant Professor, Faculty of Computer Application, AMPICS, Ganpat University, Kherva, Gujarat, India. Email: apg01@ganpatuniversity.ac.in



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ABSTRACT

Autism Spectrum Disorder (ASD) diagnosis has traditionally relied on subjective clinical observations and behavioural assessments, which have the potential for misinterpretation. The incorporation of machine learning technology into autism diagnosis represents a paradigm shift, promising higher accuracy and objectivity. This article gives a detailed comparison of autism diagnosis techniques that make use of machine learning. We illuminate the diverse landscape of machine learning applications in autism diagnosis by investigating mobile applications, neuro imaging approaches, natural language processing, and genetic data analysis. We highlight the possibility for early and accurate autism diagnosis, as well as the implications for future therapies and interventions, by examining the benefits and limitations inherent in this transformative approach. Furthermore, we look at the ethical issues that come with this technological advancement, emphasising the necessity of data protection, transparency, and patient permission. This article aims to provide healthcare professionals, researchers, and carers with insights into the expanding field of machine learning in autism diagnosis, allowing them to make more educated decisions for the benefit of people on the autistic spectrum. We find ourselves at the crossroads of innovation and compassion as we embark on this comparative adventure, altering the future of autism evaluation via the prism of machine learning technologies.

Keywords: ASD Diagnosis tools, Machine Learning, Autism, Behavioural Analysis, neuro imaging, natural language processing.





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INTRODUCTION

The merging of cutting-edge technology and machine learning algorithms has resulted in a dramatic evolution in the field of autism diagnosis in recent years. Autism spectrum disorder (ASD) is a complex neurological illness that affects people throughout their lives. Traditionally, clinical observations, standardised questionnaires, and behavioural tests were used to diagnose ASD. While these procedures have proven to be effective, they frequently lack the precision, efficiency, and objectivity required to address the numerous and subtle aspects of autism. However, the introduction of machine learning technology has ushered in a new era of diagnosis, one that promises to improve the accuracy and speed of recognising autism while lowering the subjectivity associated with older methods. This article conducts a thorough comparison of autism diagnosis methods that have harnessed the power of machine learning, exposing the astounding advances that are influencing the future of autism assessment.[1,2] As we progress through this investigation, we will examine a variety of techniques and methodologies, each of which contributes to our understanding of autism diagnosis. From mobile applications that analyse behavioural patterns to advanced neuro imaging techniques, this article provides a comprehensive look at the intersection of machine learning and healthcare, all with the goal of improving the lives of people with autism. We will investigate the benefits and limitations of machine learning in autism diagnosis, the important participants in this new environment, and the ethical questions that accompany this shift in our attempt to comprehend these revolutionary discoveries. We hope to shine light on how these technologies are set to revolutionise how we perceive, assess, and support individuals on the autistic spectrum as we traverse the convergence of cutting-edge technology and autism. [3,4, 23]

Overview of Autism Diagnosis

Autism Spectrum Disorder (ASD) diagnosis is a complex and multidimensional procedure that is critical for early identification and intervention in individuals with autism. Autism has traditionally been diagnosed primarily through clinical observation, standardised questionnaires, and behavioural evaluations. These strategies have been used by healthcare professionals such as paediatricians, psychiatrists, and psychologists to detect the typical traits and behaviours linked with autism. [5,10]

Clinical Observation

Clinical observation is a fundamental component of autism diagnosis. Through direct interactions and systematic examinations, healthcare practitioners analyse an individual's social communication abilities, repeated behaviours, and sensory sensitivity. While clinical observation gives useful insights, it is subjective and subject to vary among practitioners. [5,6]

Standardized questionnaires

In the diagnostic process, standardised questionnaires such as the Autism Diagnostic Interview-Revised (ADI-R) and the Social Communication Questionnaire (SCQ) are widely utilised. These questionnaires collect information about a child's behaviours, development, and social interactions based on carer reports. [7]

Behavioural Assessment

The Autism Diagnostic Observation Schedule (ADOS) and other behavioural examinations aim to provide a standardised framework for evaluating an individual's social and communicative skills. These exams frequently include particular tasks and activities that assist specialists in evaluating a person's behaviour and reactions. Despite their historical relevance, conventional diagnostic approaches have disadvantages such as subjectivity, potential bias, and reliance on human judgement. Furthermore, diagnosing autism in very young children is particularly difficult because they may not yet exhibit the complete range of symptoms. With the incorporation of machine learning technologies, there has been a dramatic transformation in the approach to autism diagnosis in recent years. This shift has created new opportunities for improving the accuracy, efficiency, and objectivity of autism diagnosis. The





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following sections of this article will conduct a comparative review of autism diagnosis methods that use machine learning, providing light on innovations that have the potential to reshape the landscape of autism assessment. [8]

Machine learning in autism Diagnosis

In recent years, the convergence of healthcare and technology has resulted in a game-changing synergy that has the potential to transform how we diagnose and comprehend Autism Spectrum Disorder (ASD). The fearsome capability of machine learning - an advanced area of artificial intelligence that equips computers with the ability to learn from data and generate predictions without explicit programming - is at the heart of this disruptive evolution. [2] Autism diagnosis has traditionally relied on the expertise and subjective judgement of healthcare professionals. Clinical observations, carer reports, and standardised assessments have long been used to diagnose autism, providing vital insights into the behavioural patterns and developmental milestones of people on the autism spectrum. While these methods have clearly been useful, they are not without limitations, such as subjectivity and the possibility of human mistake.[1,5] Machine learning, on the other hand, adds a new and objective component to autism diagnosis. It enables diagnostic instruments to process massive amounts of data, find nuanced patterns, and make predictions with unrivalled accuracy. It's equivalent to presenting the medical community with a precise and untiring diagnostic helper capable of quickly and consistently analysing different and sophisticated data.

Machine learning technology enables the development of specialised algorithms that can filter through complex behavioural and medical data, revealing hidden insights and patterns that would otherwise go undiscovered. These algorithms may analyse a wide range of characteristics, such as behavioural observations, genetic markers, brain imaging data, and more, to provide a complete and data-driven assessment of a person's likelihood of being on the autistic spectrum.[3,4,23] The goal of this essay is to conduct a thorough comparison of the many autism diagnosis techniques that use machine learning, shining light on the astounding breakthroughs that are transforming the landscape of autism assessment. We hope to provide a comprehensive understanding of how machine learning is poised to redefine not only how we diagnose autism, but also how we comprehend and support individuals on the autism spectrum by examining the benefits, challenges, and ethical considerations inherent in this integration of technology. As we continue on this adventure of discovery, we are on the verge of a transformative era in healthcare, driven by the marriage of innovation and compassion.[11]

Impact of Machine Learning In Autism Diagnosis

The incorporation of machine learning into the field of autism diagnosis has ushered in a new age in healthcare. Machine learning has the potential to greatly improve the precision, efficiency, and objectivity of autism diagnosis by using modern algorithms and the ability to analyse large and diverse datasets. Here, we look at how machine learning is changing the diagnostic environment for autism and the several benefits it offers.

Improved Accuracy and Objectivity

The potential for increased accuracy is one of the key benefits of adding machine learning into autism diagnosis methods. Machine learning models can analyse a large amount of behavioural, clinical, and biological data to provide objective evaluations, lowering the risk of human mistake and subjectivity.

The ability of machine learning algorithms to find nuanced patterns and relationships within datasets allows them to uncover subtle behavioural and physiological signs of autism that may defy standard diagnostic methods.

Early Detection and Intervention

Machine learning algorithms can detect indicators of autism at a younger age, allowing for timely intervention and assistance. Early diagnosis is critical for offering personalised therapy and resources to those on the autism spectrum, which can dramatically enhance their long-term outcomes. In extremely young children, machine learning technologies can detect tiny behavioural patterns that are symptomatic of autism, allowing healthcare professionals and carers to take preventative measures.





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Multimodal Data Integration

Medical records, genetic information, neuro imaging data, and natural language are just a few of the data kinds that machine learning approaches can handle. Machine learning-based techniques can provide a thorough and holistic knowledge of an individual's condition by merging these various data sources. This multimodal approach not only aids in diagnosis, but also leads to a better understanding of autism's underlying causes and heterogeneity.

Tailored and personalized Diagnosis

Individualised assessments can be provided by machine learning by taking into account a person's unique profile and traits. This personalisation ensures that each individual's diagnosis and treatment plans are specifically customised to their needs and challenges. Machine learning assists in offering more exact recommendations for therapeutic therapies by adapting to the intricacies of each instance. [2,3,4,11,23]

Enhanced Research and Insights

Machine learning methods have the ability to shed light on the complexities of autism. They can find previously unknown correlations, biomarkers, and risk factors, which help comprehend the condition and its underlying causes. Machine learning's data-driven approach fosters continued study and innovation in the field of autism, perhaps leading to breakthroughs in treatment and support. We will conduct a comparative analysis of specific autism diagnosis techniques that use machine learning. We hope to provide a thorough overview of the ever-changing environment of autism evaluation by investigating these tools, their capabilities, and their potential impact on healthcare and the lives of individuals with autism.[12,13]

Benefits of Machine Learning in Autism diagnosis

The use of machine learning technology into autism diagnosis tools provides a slew of compelling benefits that are changing how we view, diagnose, and support people on the autistic spectrum. As we compare these unique technologies, it's critical to emphasise the advantages they bring to the forefront of autism diagnosis

- 1. Machine learning algorithms can quickly and precisely analyse large datasets, minimising the margin for human error and subjective judgement in diagnosis.
 - These methods provide a higher degree of accuracy in discriminating between neurotypical individuals and those with autism by recognising minor behavioural, genetic, or neurological patterns.
- 2. Early detection is critical for prompt intervention and support, which can dramatically improve outcomes for people with autism.
 - Early indications of autism can be detected using machine learning-based methods, allowing healthcare professionals and carers to begin therapies and interventions at an earlier age.
- 3. Machine learning may integrate a wide variety of data types, such as behavioural observations, medical records, genetic information, neuro imaging data, and textual reports.
 - Given the disorder's multidimensional nature, this comprehensive approach provides for a more complete view of autism.
- 4. Machine learning algorithms provide objective and consistent evaluation, eliminating the impact of human bias in the diagnostic process. This impartiality assures that each assessment is completely based on data-driven analysis, which improves the diagnostic outcome's reliability.
- 5. Machine learning algorithms can personalise diagnoses and treatment suggestions based on each person's unique qualities and needs. Personalization ensures that interventions are particularly intended to address the person with autism's challenges and capabilities, resulting in more effective assistance.
- 6. Machine learning-powered tools promote ongoing study and innovation in autism diagnosis and treatment. These techniques contribute to a better understanding of autism's underlying causes by analysing large datasets, potentially leading to advances in therapies and interventions.





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- 7. Through digital platforms, machine learning-based technologies can be made available, allowing for remote evaluations and decreasing the pressure on healthcare systems. Because of this accessibility and efficiency, more people, including those in underserved areas, can get autism diagnostic services.
- 8. We will investigate and contrast specific autism diagnosis systems that use machine learning technologies to accomplish these advantages. Understanding the tremendous benefits of these tools allows us to see how they are set to reshape the landscape of autism assessment and provide more effective and compassionate care for people on the autistic spectrum.[2,3,4,13,23]

CHALLENGES AND LIMITATIONS

Data Quality and Integrity

Limited Datasets

For training and validation, machine learning models require vast, diverse datasets. Collecting sufficient and representative data in the context of autism can be difficult, especially for rare subtypes or specialised demographics. The varied nature of autism causes data quality problems. Variability in behaviour, symptoms, and clinical profiles might make it difficult to create robust models.

Bias and Generalization

Data Bias

Machine learning models have the potential to perpetuate bias in training data. Models may deliver erroneous or biased diagnoses if data gathering is biased or unrepresentative. It is a continuous problem to ensure that machine learning models generalise well to varied populations and demographics.

Interpretability

Black-Box Models

Some machine learning models, particularly deep learning algorithms, might be difficult to understand. Because of this lack of transparency, it might be difficult to grasp how and why a diagnosis was made, thus generating problems in clinical practise.

Regulatory and ethical concern

Data Privacy

The collecting and analysis of sensitive medical and personal data for diagnosis raises privacy and security concerns. It is vital to keep patient information secure. It is both ethical and legal to obtain informed consent for the use of patient data and the use of machine learning algorithms.

Validation and Clinical Integration

Clinical Validation

Machine learning models must be rigorously validated in order to be reliable and safe in clinical practise. This method can be time-consuming and resource-intensive. It is difficult to integrate machine learning techniques into clinical processes and ensure that they support the diagnostic process without adding additional burdens to healthcare personnel.

Interdisciplinary collaboration

Collaboration in Healthcare and Technology

Effective collaboration between healthcare practitioners and data scientists is critical. Continuous interdisciplinary communication is required to ensure that machine learning models correspond with clinical competence and patient-centred care.





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Diagnosis Complexity Autism's Complexity

Autism is a complicated and varied disorder with significant diversity in symptoms. Recognising these obstacles and limitations is critical for gaining a fair view of machine learning's involvement in autism diagnosis. These reservations do not diminish the potential of machine learning technology, but rather highlight the importance of responsible and intentional integration, ongoing study, and interdisciplinary collaboration. By tackling these issues, we can fully realise the potential of machine learning to improve autism diagnosis and care for people on the autistic spectrum.[2,3,4,12,13,14,23]

Comparison of Autism Diagnosis tools:

The incorporation of machine learning technologies into the realm of autism diagnosis has resulted in a dynamic shift in the way we approach autism spectrum disorder (ASD) assessment. The diagnostic tool landscape has grown to include a wide range of advances, each using the promise of artificial intelligence to improve precision and impartiality. In this section, we will conduct a comparative examination of some of the most popular autism diagnosis systems that have made machine learning a key component of their diagnostic process. Shown in the table[1], we will examine each of these tools in depth, examining their capabilities, performance, and possible impact on the landscape of autism diagnosis. We hope that by doing this comparative study, we will be able to provide significant insights into the options and considerations that healthcare professionals, carers, and people on the autism spectrum may encounter while looking for the most effective and trustworthy diagnostic instruments.[2,3,4,11,23] Here in the figure 1 we have show to range of sensitivity result of various autism diagnosis tools also figure 2 shows to range of specificity result of various autism diagnosis tools. The result may vary on the quality of data and age of candidate for all above mention tools.

Case study

We give a collection of case studies and examples emphasising real-world tools and their impact to provide clear insights into the practical applications of machine learning technology in autism diagnosis

ADOS2

The classic ADOS-2 assessment was supplemented with machine learning algorithms in a research study conducted at a leading autism diagnosis centre. These algorithms analysed video footage and gathered behavioural data from youngsters undergoing evaluation. The machine learning-enhanced ADOS-2 revealed a significant improvement in diagnostic accuracy. It successfully identified minor behavioural indicators that human physicians would miss, minimising false negatives and boosting overall autism diagnosis precision.[15,24]

Mobile App

To screen for autism in toddlers, researchers and technology specialists collaborated to create a smart phone application. The software analysed video records of children's interactions and play using machine learning algorithms. The app for parents and carers effectively spotted early indicators of autism. It gave risk assessments to carers, allowing them to seek expert evaluation and assistance at an early age. The tool's ease of use and accessibility were critical to its success.[16,17]

Neuro imaging and Deep learning for Brain pattern analysis

Deep learning models were applied to neuro-imaging data, including fMRI scans of children with and without autism, at a prestigious research facility. The goal was to find distinct brain activity patterns linked to autism. The deep learning approach revealed different brain activity patterns in children with autism, offering light on the disorder's neuronal foundation. These patterns were subsequently used to improve diagnostic accuracy and acquire insight into the neurological abnormalities associated with [18,19]





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NLP for textual data analysis

Researchers worked with a large healthcare institution to develop natural language processing (NLP) models that analysed text from patient reports and clinical notes. The goal was to find language markers and behavioural features linked to autism. For instance, NLP-driven research showed linguistic patterns, such as the use of specific words or phrase structures, which were associated with autism diagnosis. These linguistic cues were added into diagnostic algorithms, enhancing the diagnosis process even further. Above case studies and examples demonstrate the real-world application of machine learning technologies in autism diagnosis. They demonstrate the adaptability of machine learning technologies in a variety of domains, ranging from clinical assessments to early screening and neuroimaging analyses. By looking into these real-world applications, we obtain a better grasp of machine learning's potential for improving the accuracy, objectivity, and accessibility of autism diagnosis. In the following sections, we will continue to investigate and compare these instruments, shedding light on their importance in the field of healthcare and autism assessment. [20, 21, 22]

CONCLUSION

The landscape of autism diagnosis is experiencing a tremendous upheaval, powered by machine learning technology's extraordinary potential. This article's journey has shown the manner in which these novel tools are transforming the diagnostic procedure for Autism Spectrum Disorder (ASD). As we conclude this investigation, it is clear that the convergence of artificial intelligence and healthcare is paving the way for the future of autism diagnosis. The benefits of machine learning in autism diagnosis are numerous and varied. These instruments are superior in terms of accuracy and precision, they facilitate early identification and intervention, and they give objectivity and consistency in the diagnostic process. Machine learning algorithms provide a complete understanding of autism that outperforms traditional methods by combining data from multiple sources and employing a multidimensional approach. We met some of the important participants in this revolutionary landscape during our comparative analysis tools that analyse behavioural patterns, neuro imaging data, linguistic markers, and more. Each of these tools marks an important step towards more accurate, accessible, and personalised autism diagnosis. However, like with any revolutionary leap forward, we must be cognizant of the problems and ethical implications that this shift will entail.

Data privacy, transparency, accountability, and bias mitigation are not just nice-to-haves; they are critical foundations that sustain the ethical integrity of these diagnostic tools. Our ethical responsibility is to keep patients' rights and well-being at the forefront. We find ourselves at the crossroads of innovation and compassion as we consider the future of autism diagnosis. The machine learning technologies discussed in this article are more than just scientific advancements; they are also strong instruments of empathy, providing hope to people on the autism spectrum and their families. They represent the promise of more precise, rapid, and personalised assistance. Finally, the route forward in autism diagnosis is one distinguished by creativity, complexity, and ethical responsibility, aided by machine learning technologies. It's a trip that has the potential to change how we perceive, evaluate, and support people with autism. We can usher in a more inclusive, equitable, and empathic healthcare era with continuing interdisciplinary collaboration, adherence to regulatory standards, and a strong commitment to ethical principles. The future of autism diagnosis is here, and it appears to be brighter than ever.

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Table 1: Comparison of Machine learning based Autism screening tools]

Screening Tool Name	Age	Conducted By	Assessment Type	Length of Assessment	Reliabilit y	Validit y	Sensitivit y	Specificit y
Autism Diagnostic Observation Schedule (ADOS)	All ages	Trained clinician	Standardize d observation -based assessment	30-60 minutes	High	High	0.79-0.97	0.67-0.91
Childhood Autism Rating Scale (CARS)	2 years and older	Trained clinician	Behavior rating scale	15-30 minutes	Moderate to high	High	0.78-0.96	0.86-0.94
Social Communication Questionnaire (SCQ)	4-40 years	Parent or caregiver	Parent- reported screening tool	10-15 minutes	High	High	0.77-0.96	0.60-0.93
Social Responsiveness Scale (SRS)	4-18 years	Parent or teacher	Parent or teacher- reported screening tool	15-20 minutes	High	High	0.68-0.89	0.74-0.91

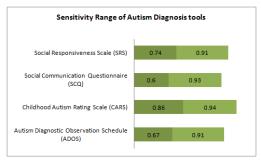


Figure 1: Sensitivity result analysis of Autism Diagnosis tools

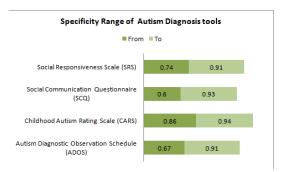


Figure 2: Specificity result analysis of Autism diagnosis tools





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RESEARCH ARTICLE

A Study on Flow of Market Information in the Supply Chain of Banana in Andhra Pradesh

Mehazabeen, A1 and G. Srinivasan2*

¹Assistant Professor, Department of Agriculture and Business Management, Centurion University of Technology and Management, Odisha, India.

²Associate Professor, Department of Agri. Economics, Annamalai University, Tamil Nadu, India.

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*Address for Correspondence

G. Srinivasan

Associate Professor, Department of Agri. Economics, Annamalai University, Tamil Nadu, India.



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ABSTRACT

Banana (Musa paradisica L.) is a vital cultivated herbaceous food crop on a global scale, with significant export potential. Banana cultivation in the Andhra Pradesh state has expanded to 1.12 lakh hectares, with a production of 63.84 lakh MT in 2019. Y.S.R. district leads the state in production, contributing 8.67 lakh MT. The study was undertaken with the objective of (i)To analyse the flow of marketing information among farmers within the supply chain of banana in Andhra Pradesh (ii) To analyse flow of marketing information among the banana traders and policy suggestions. For this study, a multistage stratified random sampling method was used. The data were analyzed and interpreted in tabular form and analyzed data were expressed in averages and percentages for the better understanding. The results from the study showed that farmers who were aware of the area of banana cultivation in the current season and previous season as well as prices were very meagre (7.5 per cent) in the study area. Hence, the study suggested that market information about cultivation area, arrivals and prices of banana should be provided by convenient of market stakeholders through ICT. The study also suggested for the establishment of institutional marketing facilities, including regulated and cooperative market, equipped with proper infrastructure to encourage banana farmers in the study area.

Keywords: Marketing intermediaries, Market information, Institutional Marketing





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INTRODUCTION

Cultivating horticultural crops plays a fundamental role in enhancing our nation's prosperity, as it directly influences the well-being and economic fortune of its people while also promoting their overall health(Vision, 2050 IIHR, 2014). Banana (Musa paradisica L.) stands as a crucial cultivated herbaceous food crop globally, boasting substantial export potential. Employing high-tech methods for its cultivation proves economically sound, resulting in enhanced productivity, superior product quality, and early crop maturity. Such produce commands a premium price in the market. Marketing information helps the banana farmers in making decisions inmarketing their produce. The farmers cannot know the prices prevailing in the markets, as the market committees can disseminate information only in respect of their own markets. Therefore, farmers are left with no alternative but to dispose of their products in the nearest market, even at uneconomic prices. Market information is equally needed by other market participants in arriving at optimal trading decisions. At this juncture, the study was undertaken with the objective of (i)To analyse the flow of marketing information among farmers within the supply chain of banana in Andhra Pradesh (ii)To analyse flow of marketing information among the banana traders and policy suggestions.

Design of the study

The Y.S.R. district in Andhra Pradesh was purposively chosen from among the 13 districts due to its leading position in both banana cultivation area covering 16,731 hectares, and production8,67,663 metric tons. For this study, a multistage stratified random sampling method was used, with Andhra Pradesh state as the overall population. Districts were chosen as the first-stage units, Mandals within the districts as the second-stage units, villages within the blocks as the third-stage units, and ultimately, individual farmers served as the final sampling units.16 villages which are above the mean area of banana cultivation of the district, which consists of nine villages from Lingala Mandal and seven villages from Vemula Mandal were selected for this study. A sample of 320 farmers was randomly selected using the probability proportionate method, accounting random numbers for identification. Data collection involved surveying a sample of 120 market intermediaries, including commission agents, pre-harvest contractors, wholesalers, and retailers, engaged in banana marketing at the Pulivendula market.

Tools of Analysis
Descriptive Statistics
Simple Tabular Analysis
The data were analyzed and interpreted in tabular form.
Averages and Percentages

The analyzed data were expressed in averages and percentages for the better understanding.

RESULTS AND DISCUSSIONS

Flow of Market Information in the Supply Chain of Banana Farmers Awareness on Market Information

The details of the farmer's awareness on the market information are given in Table 1. The table indicates that the majority of the farmers that is 86.25 per cent are aware of prices of banana in the local market while 70.62 per cent of the farmers are aware of the arrivals of banana in the local market. About 32.5 per cent of the farmers are aware of the quality and grades of the banana and 38.75 per cent of the farmers are aware of banana prices in other markets. Only 7.5 per cent of the farmers are aware of the area of crop sown in the state.

Traders awareness on market information

The details of the trader's awareness on various types of market information are represented in the Table2. The table revealed that all the traders are aware of the prices of banana in the local market. About 97.50 per cent of the traders are aware of arrivals in the local market and 85.83 per cent of the traders are aware of prices in the reference markets





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like Gorakhpur, Banglore, Delhi, Hyderabad, Tirupati, Hubli, Dharwad, and Haveri markets. About 90.83 per cent of the traders are aware of arrivals of banana in the reference market. About 73.33 per cent of the traders are aware of the area under the crop of banana and about 63.33 per cent of the traders are aware of the production of banana in the state. Only 20.00 per cent of the traders are aware of the export or import of banana from the region.

Farmer's Source of Market Information at Village Level and Market Level

The details source of market information to the farmers is presented in Table 3. Themajor share forthe source of market information is by television in agriculture-related programmes like Pasidipantalu, Kisan, Annadaata and other programmes (87.5 per cent) followed by the newspaper (48.75 per cent) and radio programmes like Kissan Vani (14.68 per cent) and only 12.18 per cent of the farmers are using the internet as a source of market information. About 49.37 per cent of the farmers are aware of market information through the state horticultural department. About 77.5 per cent of the market information is through commission agents and 73.75 per cent through traders and 23.75 per cent through input dealers.

Source of Market Information of Traders

The details of source of market information for the traders are presented in Table 4. The table clearly indicates that the 93.33 per cent of the traders are availing the market information through the contacts in other markets over the phone, whereas, 85.00 per cent of the traders are aware of market information through fellow traders followed by newspapers (60.00 per cent), market information cell, (61.67 per cent), internet (56.67 per cent) and magazines (34.16 per cent).

CONCLUSION

In banana market, farmers who were aware of the area of banana cultivation in the current season and previous season as well as prices were very meagre (7.5 per cent) in the study area. Hence, the study suggested that market information about cultivation area, arrivals and prices of banana should be provided by convenient of market stakeholders through ICT. The study also suggested for the establishment of institutional marketing facilities, including regulated and cooperative market, equipped with proper infrastructure to encourage banana farmers in the study area.

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Table 1. Farmer's Awareness on Market Information

Sl. No.	Type of Market Information	No. of Respondents	Percentage
1	Prices in the local market	276	86.25
2	Arrivals in the local market	226	70.62
3	Quality and grade of the produce	104	32.5





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4 Prices in the other market 124 38.75 5 Area of crop sown in the state 20 7.5

(n=320) Note: The percentage do not add up to 100 due to multiple or no response

Table 2. Trader's Awareness on Market Information

Sl. No.	Type of Market Information	No. of Respondents	Percentage
1	Arrivals in local market	117	97.50
2	Arrivals in reference market	109	90.83
3	Prices in local market	120	100.00
4	Prices in reference market	103	85.83
5	Area under crop	88	73.33
6	Production of banana in the state	76	63.33
7	Quality/Grade required	92	76.67
8	Export and import	24	20.00

(n=120) Note: The percentage do not add up to 100 due to multiple or no response.

Table 3.Farmer's Source of Market Information at Village Level and Market Level

Sl. No.	Source of Market Information	No. of Respondents	Percentage
1	Radio	47	14.68
2	Newspaper	156	48.75
3	Television	280	87.50
4	Internet	39	12.18
5	Friends/Relatives/Neighbours	124	38.75
6	State horticulture department	158	49.37
7	Traders	236	73.75
8	Commission agents	248	77.50
9	Input dealers	76	23.75

(n=320) Note: The percentage do not add up to 100 due to multiple or no response.

Table 4. Source of Market Information of Traders

Sl. No.	Source of Market Information	No. of Respondents	Percentage
1	Radio	21	17.50
2	Television	54	45.00
3	Newspaper	72	60.00
4	Magazines	41	34.16
5	Internet	68	56.67
6	Contact in other markets over phone	112	93.33
7	Fellow traders	102	85.00
8	Govt. publications	24	20.00
9	Market information cell	74	61.67

(n=120) Note: The percentage do not add up to 100 due to multiple or no response.





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RESEARCH ARTICLE

Unveiling Scholarly Insights: Analyzing Social Media Advertising Literature through Bibliometric Analysis, Science Mapping **Visualization Techniques**

Nitika* and Mohit

Research Scholar, Institute of Management Studies and Research, Maharshi Dayanand University, Rohtak, Haryana, India.

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*Address for Correspondence

Nitika

Research Scholar, Institute of Management Studies and Research, Maharshi Dayanand University, Rohtak, Haryana, India. Email: nitikamalik.rs.imsar@mdurohtak.ac.in



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ABSTRACT

This expansive study undertakes a thorough exploration of the social media advertising (SMA) landscape from 2014 to 2023, employing advanced bibliometric analysis tools such as VOS viewer, Bilioshiny, and SPSS software. A meticulous examination of 242 documents from 137 diverse sources, including journals and books, paints a comprehensive picture of SMA research. The study unfolds with notable characteristics, including a robust 16.18% annual growth rate, showcasing the field's increasing influence and relevance. The relatively low average age of 2.96 years of the documents indicates a commitment to incorporating the latest research in the rapidly evolving domain. Each document's average of 20.24 citations underscores the scholarly impact of the study, while the extensive use of 14,915 references attests to the depth and thoroughness of the research. Enriched document contents, comprising 390 Keywords Plus and 784 Author's Keywords, contribute to a nuanced understanding of the subject matter. Collaboration emerges as a hallmark of this research, involving 630 authors, 26 of whom authored single-authored documents. The collaborative spirit extends globally, with an average of 2.94 co-authors per document, and 26.45% of collaborations being international, demonstrating a truly global perspective. Shifting focus to the SMA landscape, the study's robust Scopus database search protocol reveals a 16.18% annual growth rate, peaking at 198 publications in 2022. The United States, India, and China emerged as major contributors, highlighting the global recognition of SMA. Country collaboration analysis identifies extensive partnerships, particularly with Korea and China. Prolific authors and influential figures in co-citation analysis, including the highly cited work by Lee, Hosanagar, and Nair on advertising content's impact on user engagement, provide a comprehensive view of the SMA research landscape. Journal rankings position the Journal of Research in Interactive Marketing as





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the most impactful source, shaping the trajectory of SMA research. Marketers, policymakers, and researchers stand to benefit significantly from the insights offered by this study, gaining valuable perspectives for enhanced campaign effectiveness, regulatory guidance, and future research directions. By synthesizing diverse themes, this study not only captures the current SMA landscape comprehensively but also steers future research by identifying emerging trends and unexplored territories in the dynamic advertising arena.

Keywords: Literature Review, Bibliometric Analysis, SMA Analysis, Co-Citations Analysis, VOSviewer, Science Mapping.

INTRODUCTION

In the digital age, the pervasive influence of social media platforms has revolutionized the way businesses and brands engage with their target audiences. One of the most impactful manifestations of this transformation is the realm of social media advertising (Cheung et al., 2023). As users continue to flock to platforms such as Facebook, Instagram, Twitter, and LinkedIn, advertisers have recognized the unparalleled potential of these spaces for reaching, engaging, and persuading consumers (McInnis et al., 2022). The dynamic and rapidly evolving nature of social media advertising has prompted researchers to delve into its multifaceted dimensions, contributing to an expanding body of knowledge in the field (Ertemel and Ammoura, 2016). This bibliometric research paper embarks on an exploration of the extensive landscape of scholarly literature related to social media advertising. Bibliometrics, a quantitative method used to analyze patterns of publication and citation within the academic literature, offers a unique lens through which to observe the evolution of research trends, seminal works, and influential authors in this domain (Donthu et al., 2021). By employing bibliometric techniques, we aim to uncover the intellectual structure, growth patterns, and emerging themes that characterize the field of social media advertising research.

The importance of this study lies not only in its potential to provide a comprehensive overview of the existing research landscape but also in its ability to guide future scholars, practitioners, and policymakers. Through a systematic analysis of publication trends, citation patterns, and interdisciplinary collaborations, we intend to identify knowledge gaps, highlight seminal contributions, and project the potential trajectories of social media advertising research. This paper serves as a valuable resource for individuals seeking to comprehend the evolution of thought within this dynamic field and make informed decisions that align with current scholarly discourse. The subsequent sections of this paper are organized as follows: Section 1 includes an introduction and background of the domain, and Section 2 introduces the methodology employed, search protocol for data extraction, selection criteria, and analytical tools used in the bibliometric analysis. Section 3 presents the key findings derived from the analysis, highlighting influential works, prolific authors, and prevalent themes within social media advertising. In Section 4, we discuss the implications of these findings, addressing potential future research directions and the practical significance of the identified trends.

BACKGROUND OF THE STUDY

The emergence of social media platforms in the early 2000s paved the way for a new era of digital advertising. Initially, platforms like MySpace and Friendster experimented with ad models, but these efforts were limited in targeting capabilities and lacked advanced analytics (Jones et al., 2008). However, the introduction of Facebook advertising in 2007 revolutionized the field (McInnis et al., 2022). Facebook's vast user base and abundant demographic data provided advertisers with unprecedented targeting opportunities (Wilson et al., 2012). They could reach specific demographics, interests, and behaviors using engaging ad formats like photos and videos (Nadkarni and Hofmann, 2012). Facebook continuously enhanced its advertising platform with features such as custom audiences and dynamic advertising, enabling companies to optimize their campaigns and achieve higher returns on





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investment (Hansson et al., 2013). This success led other social media platforms like Instagram, Twitter, LinkedIn, and Pinterest to enter the advertising space, each developing its unique ad formats and targeting capabilities to cater to different demographics and user preferences (Vishwanath et al., 2009). Brands recognized the advertising potential on these platforms, allowing them to effectively engage with their target audience. As social media platforms grew, so did research in the field of social media advertising. The first study, conducted in 1996 by Pack and Thomas, explored whether net casting would shape the future of radio or remain a technological novelty. This study laid the groundwork for subsequent research in this domain. Chawla et al. conducted a study in 1999 to understand consumer sentiments toward internet usage. Sheehan and Hoy investigated consumer attitudes and perceptions regarding privacy and online advertising practices. Over time, numerous studies have contributed to the understanding of social media advertising. In the present context, Lee et al.'s study published in 2018 has received the highest number of citations. This study examined the relationship between consumer engagement and advertising effectiveness, providing valuable insights into the field.

OBJECTIVES OF THE STUDY

The objectives of this bibliometric analysis encompass a comprehensive exploration of the social media advertising (SMA) domain from 2014 to 2023. Firstly, the temporal pattern of annual publications will be scrutinized to discern evolving trends and shifts in research focus over the specified timeframe. Secondly, the analysis aims to identify the most productive countries in the SMA domain, shedding light on global contributions to the field. Thirdly, the collaboration pattern among countries will be examined, providing insights into international research networks within SMA. Moreover, a detailed examination of the most productive authors, universities, and sources in SMA analysis will be conducted to recognize influential contributors and authoritative publications. Additionally, the cocitation pattern within the domain will be explored, highlighting interconnected works that have significantly influenced SMA research. The analysis will also delve into the examination of the most cited publication in the SMA domain, elucidating key foundational works. Furthermore, the study will employ a three-field plot analysis to provide a visual representation of the domain's intellectual structure. Additionally, the co-occurrence of keywords, trend topics, and the thematic map of the domain will be analyzed to unveil prevalent themes and emerging areas of interest. Finally, the research aims to identify major themes on which research work is conducted in the SMA domain, offering a holistic understanding of the current landscape and potential directions for future exploration.

MATERIALS AND TECHNIQUES

Bibliometric analysis, recognized as a quantitative method for assessing scientific publications, authors, and research trends (Koseoglu et al., 2016, Chaudhry & Kumar, 2023, Singh et al., 2023), serves as the cornerstone of our study's methodology in analyzing the literature on social media advertising. Embracing this approach, we leverage bibliometric tools known for their transparency and objectivity, encompassing the evaluation of citation and publication data, co-authorship, and thematic analyses (Thanuskodi, 2010). This comprehensive analysis employs performance assessment, impact measurement, and science mapping, including visualization techniques, to gain a nuanced understanding of the dynamic landscape within the field. Previous research has demonstrated the efficacy of this approach in uncovering growth trends, author keywords, and prominent journals (Andres, 2009; Subramanyam, 1983; Patra et al., 2006). Our unique contribution lies in presenting a systematic methodology that integrates Biblioshiny and VOSviewer tools. Biblioshiny facilitates efficient data retrieval and analysis, while VOS viewer constructs visually intuitive bibliometric networks (Rusydiana, 2021). Additionally, MS Excel and SPSS are instrumental in providing more in-depth insights into the domain. By employing this comprehensive toolkit, our research strives to enrich the understanding of the scholarly landscape of social media advertising, emphasizing transparency, objectivity, and an innovative methodology for uncovering key insights in this dynamic field.





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Search Protocol

The bibliographic information was gathered using the Scopus database because it has more publications than other databases like "Web of Science" and is, therefore, one of the biggest databases of journals. The Scopus database has been used for bibliometric analyses in numerous earlier works. The search string was used in the title-abstract-keyword field of Scopus. The search string used in the analysis is depicted and defined below: A variety of search procedures were utilized to ensure the coverage of all articles and documents connected to social media advertising analysis, based on the evaluation of prior literature and recommendations from academicians. The keyword "social media advertising" was used for the extraction and analysis of available literature. The Scopus database's Title-Abstract-Keywords field was searched using the mentioned keyword. Lastly, our database included journal articles only, and the data were extracted for the period of 10 years (2014 to 2023). Additionally, the subjects were limited to business management, accounting, psychology, multidisciplinary, and arts and humanities. Moreover, the analysis only considered manuscripts that were written in English.

DATA ANALYSIS AND FINDINGS

This figure above provides a comprehensive overview of the research study conducted over the time span from 2014 to 2023, encompassing 242 documents from 137 different sources, including journals and books. The study exhibits a robust annual growth rate of 16.18%, indicative of its expanding influence and relevance over the years. The document's average age is relatively low at 2.96 years, suggesting the inclusion of recent research. Each document receives an average of 20.24 citations, indicating the scholarly impact of the study. The extensive use of references, totaling 14,915, further attests to the thoroughness and depth of the research. The document contents are enriched with 390 Keywords Plus and 784 Author's Keywords, providing a comprehensive understanding of the subject matter. The collaborative nature of the research is reflected in the involvement of 630 authors, with 26 singleauthored documents. The collaboration extends to an average of 2.94 co-authors per document, and 26.45% of these collaborations are international, showcasing a global perspective. The predominant document type is an article, comprising all 242 documents, emphasizing the study's focus on in-depth analysis and scholarly discourse. Overall, this table paints a picture of a dynamic, well-referenced, and globally collaborative research endeavor with a significant and growing impact on its field. A total of 242 documents are analyzed for the study. The trend depicted by the publications is demonstrated above. Analysis revealed that the year 2022 witnessed the maximum publication in the domain with 198 documents. The domain is still in the infant stage and growing continuously year by year. The publications have an upward trend except for the year 2020 when the documents are in a downward trend and were lesser than the general trend. In the year 2023, there are fewer studies from general trends but the reason is that data is extracted till July 2023 only, not for the whole year. The above table ranks countries based on their research output and citation impact within a specified domain.

The United States leads with 90 documents and 1740 citations, signifying a substantial contribution to the field. It is classified under North America and categorized as a developed nation. India follows as the second-ranking country with 28 documents and 303 citations, representing a significant research presence in Asia. India is classified as a developing nation, highlighting its ongoing efforts in scientific and scholarly endeavors. China, with 20 documents and 147 citations, also emerges prominently in the rankings. Like India, China is classified as a developing country, emphasizing its growing influence in the global research landscape. Australia occupies the fourth position with 16 documents and 212 citations, representing a developed nation in Oceania. This suggests a noteworthy research output and impact on the Australian scientific community. In summary, this table provides a snapshot of the research landscape in different countries, reflecting variations in research productivity and impact. It underscores the global distribution of research contributions and highlights the research strengths of both developed and developing nations. Country collaboration is analyzed in this section of the analysis. The results demonstrated that the USA collaborated in the most publications with Korea, with a total of 8 publications. The USA ranks second in collaboration with China, with a total of 7 publications. Germany and the Netherlands are ranked next, with collaboration in 3 publications each. The USA also published the same number of publications in collaboration with





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Australia, Germany, New Zealand, and the United Kingdom. Furthermore, Korea and New Zealand collaborated on 3 publications as well. The countries' collaboration map is depicted below: Single-authored and multiple-authored collaborations of countries are analyzed in this section. The USA tops the list with 57 publications, 47 of which are single-authored and 10 are multiple-authored publications. China is in 2nd position with a total of 21 publications, comprising 10 single-authored publications and 11 collaborations with multiple authors. India occupies the 3rd position with 16 publications, all of which are single-authored; no multiple-authored collaborations were done by India. After that, it is followed by Korea with a total of 11 publications, 5 of which are single-authored and 6 are multiple-authored. This table presents a ranking of authors based on the number of documents and the total citations received. The authors listed have contributed to the field of social media advertising (SMA), and their impact is measured by the number of documents they have authored and the citations those documents have received. Rana M, Arora N, and Arora T, Agarwal B emerge as the most prolific authors, with multiple documents attributed to their names. However, Arora T, and Agarwal B hold the second position but have a notably higher citation count, indicating the influential impact of their work. Boateng H, Okoe A.F. and Natarajan T, Balakrishnan J, Balasubramanian S.A, and Manickavasagam J also feature prominently, contributing to two documents each, with varying citation counts The diverse range of authors in the table signifies the collaborative nature of SMA research, with some authors co-authoring documents.

Noteworthy is the significant citation count associated with certain authors, such as Alalwan A.A. with 278 citations, indicating the substantial impact and recognition of their work within the SMA domain. Overall, this table provides insights into the productivity and impact of various authors in the field of SMA, offering a snapshot of the research landscape and highlighting key contributors and influential works. Table 7 presents the top authors in this bibliographic review, ranked by their citations per publication. The table provides the names of the authors, the number of papers published, and the total number of citations for their works. At the top of the list in the first row are authors "Lee D., Hosanagar K., and Nair H.S." closely followed in second place by Voorveld H.A.M., Van Noort G., Muntinga D.G., and Bronner F., with 321 citations for their single publication. In the third position are Lee J. and Hong I.B., receiving 302 citations for their one publication. The fourth place is secured by Alalwan A.A. with 278 citations in one publication, while Zhu Y.-Q. and Chen H.-G. rank fifth with 206 citations for their one publication. The subsequent entries showcase different authors, each with a single publication and separate citation counts. These authors have made significant contributions to the study through their individual works, emphasizing the impact of their research. Overall, Table 7 highlights the presentation of influential authors in this literature review based on their citations, shedding light on the impact and quality of their individual research contributions. When two texts often cite two different articles by two different authors, co-citation analysis is used. The top author's co-citations analysis showed that Hair J.F. topped the list with 279 citations and the strongest co-citation link strength (7736).

Moreover, Law R., with 252 citations, Sarstedt M., with 230 citations, Ringle C.M., with 222 citations, and Wang Y., with 196 citations, were among the top 5 in the list according to citations. This table provides a glimpse into the landscape of social media advertising research by ranking the most cited publications in the field. Topping the list is the work by Lee, Hosanagar, and Nair (2018), titled "Advertising Content and Social Media: Evidence from Facebook," published in the journal "Management of Science," with an impressive 399 citations. Following closely is the paper by Voorveld, van Noort, Muntinga, and Bronner (2018) from the "Journal of Advertising," exploring engagement with social media and advertising across different platforms, garnering 321 citations. The third-ranked paper by Lee and Hong (2016) investigates the predictors of positive user responses to social media advertising, emphasizing emotional appeal, informativeness, and creativity, accumulating 302 citations. Other notable contributions include Alalwan's (2018) exploration of the impact of social media advertising features on customer purchase intention, and Zhu and Chen's (2015) examination of the relationship between social media and human need satisfaction. This compilation not only underscores the diverse aspects covered in social media advertising research but also highlights the scholarly impact of these works, offering a valuable resource for academics, marketers, and researchers seeking a deeper understanding of this evolving field. The presented table offers a comprehensive ranking of sources in the realm of social media advertising, assessing their impact based on the number of documents published, total citations received, and the average number of citations per publication. At the





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forefront, the "Journal of Research in Interactive Marketing" claims the top position with an impressive tally of 16 documents and a substantial 461 citations, resulting in an exceptional average of 28.81 citations per publication. Following closely is the "International Journal of Internet Marketing and Advertising," contributing 9 documents with 28 citations and an average of 3.11 citations per publication. The "Journal of Interactive Advertising" secures the third spot, boasting 8 documents and 81 citations, yielding an average of 10.125 citations per publication. The fourth entry, "Computers in Human Behavior," stands out with 6 documents but an outstanding 531 citations, resulting in a remarkable average of 88.5 citations per publication. This ranking enables a nuanced understanding of the influence and recognition of each source within the academic discourse of interactive marketing and advertising, providing valuable insights for researchers, practitioners, and scholars in the field. This table tracks the annual publication counts for five prominent journals - "Journal of Research in Interactive Marketing," "International Journal of Internet Marketing and Advertising," "Journal of Interactive Advertising," "Computers in Human Behavior," and "Frontiers in Psychology" - from 2014 to 2023. The data provides valuable insights into the evolving research landscape within the field. In 2014, the "International Journal of Internet Marketing and Advertising" led with a single publication, followed by an increasing trend over subsequent years. "Journal of Research in Interactive Marketing" commenced its publications in 2015 and exhibited steady growth, reaching 16 publications in 2023. "Journal of Interactive Advertising" and "Computers in Human Behavior" started with zero publications but experienced gradual increases, particularly in 2018 and beyond. "Frontiers in Psychology" entered the scene in 2016 and maintained a moderate but consistent publication output. Throughout the years, 2021 witnessed a notable surge in publications across all journals, indicating a heightened focus on research in social media advertising.

The year 2022 marked a significant increase in publications for all journals, showcasing a growing interest and scholarly engagement. The data suggests a dynamic landscape with shifting research priorities, emphasizing the importance of these journals in contributing to the academic discourse on interactive marketing, internet marketing, advertising, human behavior, and psychology within the specified timeframe. Researchers and stakeholders can leverage this information to gauge the trajectory of research interest and identify potential trends within the field over the past decade. The table presents a ranking of organizations based on their scholarly output and citation impact. The "Department of Marketing at Cape Peninsula University of Technology in Cape Town, South Africa," secures the top position with 2 documents and a notable citation count of 177. Similarly, the "Department of Marketing, University of Professional Studies in Accra, Ghana," also records 2 documents but slightly fewer citations at 173. The third-ranking organization is the "Department of Retail, Hospitality, and Tourism Management" at the University of Tennessee, Knoxville, United States, with 2 documents and 59 citations. The ranking reflects the research productivity and influence of these academic departments in the field of marketing and related disciplines. The organizations from South Africa and Ghana stand out in terms of both document output and citation impact, showcasing their contribution to the scholarly landscape. The Sankey diagram in the figure illustrates the main components of three fields-authors, keywords, and journals, and their interactions. Arrow width is inversely related to flow velocity. Three local names denote the most popular authors, keywords, and sources. A chart of colored triangles indicates the main components, with relationships summarized in terms of rectangle height. The number of combinations for each unit depends on the rectangle size. The current study identified research topics and main sources, revealing "Consumer Heterogeneity and Paid Search Effectiveness:

A Large-Scale Field Experiment Econometrics" by Blake T. et al. as the most important node with 2 flow counts. The second-highest related node is "When Does Retargeting Work? Information Specificity in Online Advertising" with 1 flow count, published in the Journal of Marketing Research by Lambrecht A. and Tucker C., followed by "Control and Peripheral Routes to Advertising Effectiveness: The Moderating Role of Involvement" in the Journal of Consumer Research by Petty R.E. The provided table presents an analysis of frequently occurring keywords within a specific context. The most prominent keyword, "Social Media," appears 88 times, highlighting the central role of social platforms in the subject matter. "Social Media Advertising" closely follows with 72 occurrences, indicating a strong focus on using these platforms for promotional activities. The more general term "Advertising" is also significant, occurring 46 times, suggesting a broader exploration of promotional strategies beyond just social media. "Marketing" and "Purchase Intention" round out the list with 25 and 24 occurrences, respectively, reflecting a





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continued interest in core marketing principles and the study of consumer behavior related to the discussed topic. This keyword analysis underscores the emphasis on social media and its intersection with advertising and marketing concepts in the context of the subject being studied. The figure below depicts the graph of the same keywords built using VOSviewer. The density visualization, which reveals research hotspots, indicates that "Social Media Advertising," "Social Media," and "Advertising" are key research hotspots in the domain. The figure depicting this is presented above. The research hotspots in different years were analyzed using the Biblioshiny software, and the results demonstrated that in 2022, the terms "online advertising" with 18 occurrences, "advertising" with 10 occurrences, and "forecasting" with 7 occurrences were the most frequent topics.

Meanwhile, in 2021, "social media" with 51 occurrences, "electronic commerce" with 14 occurrences, and "consumption behavior" with 8 occurrences were the most researched topics in the domain. In 2020, this list comprised "marketing" with 85 occurrences, "advertising" with 56 frequencies, and "female" with 42 occurrences, representing the hotspots of the stated research domain area. In this section, a thematic map of the domain is created. Layers are assembled to generate thematic maps, and the data from each layer are then plotted in one or more aesthetics. Thematic evolutionary analysis is employed to identify relationships between historical patterns and evolutionary processes. The thematic analysis utilizes an inclusion index weighted by incoming terms, with a minimum cluster frequency of 5 and a minimum weighting index of 0.1. The strength of the structures that develop over time is depicted in the theme analysis map, where each node's size corresponds to the topic's total number of keywords. The top left quadrant, characterized by high density but low centrality, presents specific and underrepresented issues still undergoing major growth, themes such as "social media platforms," "commerce," "consumer behavior," "psychology," "COVID-19," and "epidemiology" fall into this niche category within the domain. Additionally, the domain exhibits declining themes like "behavioral research," "social influence," and "empirical analysis." The lower right quadrant contains basic themes with high centrality and density, including "social media advertising," "consumption behavior," "marketing," "social networking (online)," and "advertising." Motor themes in the domain area encompass "human," "article," "female," "purchase intention," "purchasing," and "sales."

MAJOR RESEARCH THEMES IN SOCIAL MEDIA ADVERTISEMENT DOMAIN

The landscape of advertising has significantly transformed with the rise of social media platforms, leading to a surge in research efforts to understand the intricate dynamics between social media advertising and various aspects of consumer behavior, effects, influencers, trust, culture, brand perception, ethics, health impact, and advertising effectiveness. This analysis synthesizes the research contributions of various authors in our domain area and also analyzes the various themes on which research work is done in the domain area of social media advertising. The stated analysis can be very crucial for future researchers and can guide them in further studies. The major themes of the domain area are stated below:

Social Media Advertising and Consumer Behavior

The influence of social media advertising on consumer behavior is a central theme. Chen & Chiu (2021) explore the impact of personalized advertisements on consumer responses. Sarılgan et al. (2022) delve into the role of emotional appeals in driving purchase intention. Asiedu (2017) investigates the impact of social media advertising on brand loyalty. This theme is also addressed by Agil et al. (2022), Sreejesh et al. (2020), Arora et al. (2020), Tian & Li (2022), and Tan et al. (2021).

Effects of Social Media Advertising

The effects of social media advertising are diverse and multi-faceted. Studies by Supotthamjaree & Srinaruewan (2021), Li et al. (2020), and Sreejesh et al. (2020) emphasize the impact on consumer attitudes and behaviors. Lee & Hong (2016) investigate the influence of social media advertising on brand awareness. Alalwan (2018) examines the effects on purchase intention. Lim & Childs (2020) study the impact on engagement, and Raji et al. (2019) investigate the role of source credibility.





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Influencers and Brand Endorsement using Social Media Advertising

Influencers and brand endorsement have become integral to social media advertising. Kim et al. (2023) focus on the effectiveness of influencer marketing. Tian & Li (2022) explore the role of celebrity endorsements in shaping consumer attitudes. Hwang & Zhang (2018) examine the impact of different types of influencers. Chiu & Ho (2023) discuss the influence of authenticity in brand endorsements. Ahmad et al. (2019) study the credibility of influencer content.

Privacy and Trust in Social Media Advertising

Privacy and trust are crucial considerations in the digital era. Fletcher (2001) explores the relationship between online privacy concerns and information disclosure. Bright et al. (2022) investigate the impact of privacy concerns on ad effectiveness. Efendioglu & Durmaz (2022) examine the role of trust in shaping consumer responses. Alkis & Kose (2022) delve into the impact of privacy-related messages on trustworthiness.

Cultural and Demographic Influences in Social Media Advertising

Cultural and demographic factors play a significant role in social media advertising effectiveness. Nguyen et al. (2023) analyze the influence of culture on consumer responses. Bui (2022) examines the role of cultural dimensions in shaping perceptions. Muk et al. (2014) explore the influence of demographics on advertising effectiveness. Thornhill et al. (2017) study the impact of cultural values on consumer attitudes, and Mansour (2015) investigate the influence of cultural norms.

Brand Perception and Consumer Engagement using Social Media Advertising

Brand perception and consumer engagement are pivotal for successful advertising. Chan (2022) examines the impact of brand personality on consumer engagement. Yang et al. (2021) explore the role of consumer-brand identification. Cheung et al. (2022) investigate the influence of emotional appeals on brand perception. Adamopoulos et al. (2018) study the effects of user-generated content on consumer engagement. Jaitly & Gautam (2021) explore the relationship between brand loyalty and consumer engagement.

Ethical and Psychological Aspectsof Social Media Advertising

Ethical and psychological aspects of social media advertising are essential considerations. Paramita & Septianto (2021) study the impact of ethical considerations on consumer responses. Bui et al. (2021) examine the role of ethical appeals in advertising effectiveness. Kim et al. (2023) focus on the psychological effects of social media advertising. Chauhan & Shukla (2016) explore the influence of psychological factors on ad effectiveness. Feng & Xie (2018) analyze the impact of psychological factors on consumer behavior.

Health-related and Societal Impact

The health-related and societal impacts of social media advertising are emerging areas of research. Al-Jabir et al. (2020) discuss the role of social media advertising in promoting public health. Rambe& Jafeta (2017) examine the impact of health-related advertising on consumer behavior. Athey et al. (2023) focus on the societal consequences of health-related advertising. Seberíni & Tokovska (2021) study the role of social media advertising in shaping health-related behaviors. Coulthard et al. (2021) analyze the effects of social media advertising on societal norms.

Social Media Advertising Effectiveness and Engagement

Advertising effectiveness and engagement are core concerns in the domain. Lou & Koh (2018) explore the factors influencing ad effectiveness. Dankwa (2021) investigates the role of engagement in enhancing ad effectiveness. Rizomyliotis et al. (2021) study the relationship between ad content and consumer engagement. Lee & Hong (2016) focus on the effects of social media advertising on ad recall and engagement. In conclusion, social media advertising is a multifaceted field encompassing various dimensions of consumer behavior, effects, influencers, trust, culture, brand perception, ethics, health impact, and advertising effectiveness. The research contributions of the mentioned authors shed light on the intricate interactions within these themes, contributing to a comprehensive understanding





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of the evolving landscape of social media advertising. These research themes also can be employed by future researchers for further investigation in the domain area.

DISCUSSION

Social Media Advertising (SMA) analysis is an area of study that's gaining traction in the academic world day by day. Our study adds to what already exists in the domain of SMA analysis by using two approaches: bibliometric analysis and text mining approach. These methods help us get a more complete understanding of SMA research and help us identify the important research journals, institutions, authors, and trends in the field. What we found is that SMA analysis is still in its early stages in academia. It's clear from the analysis that the research on SMA is increasing, especially since 2017. We noticed that researchers started collaborating more after 2014, with a peak in 2018. From 2016 to 2020, several influential studies were published by authors like "Lee D., Hosanagar K., Nair H.S. (2018), Voorveld H.A.M., Van Noort G., Muntinga D.G., Bronner F. (2018), Lee J., Hong I.B. (2016), and Alalwan A. and others. These works have shaped how we understand SMA analysis. Geographically, most SMA analysis publications come from the United States, followed by India and China. This shows that research efforts are spread across the world. Authors like ALALWAN A.A. have received the most citations, and Rana M. and Arora N. have contributed significantly in terms of the number of publications.

Their work, along with authors like Lee D., Hosanagar K., Nair H.S., has had a strong impact on the academic discussions in this field. This suggests that SMA analysis is interdisciplinary and involves various perspectives. The Department of Marketing, Cape Peninsula University of Technology, Cape Town, South Africa, stands out for having a high number of SMA analysis publications, showing their commitment to this area of research. In addition, our study noticed that new technologies are being integrated into SMA to make it more strategic and data-driven. This shift is helping to address issues related to people and audiences. Social media advertising software is also making it easier for businesses to advertise on platforms like Facebook and Twitter, reaching millions of people. However, there's still a gap in research, especially in the context of India. In conclusion, our study gives us a better understanding of SMA analysis, its influencers, and how it's evolving. By using different analysis methods, we've been able to see the bigger picture of this dynamic field. We've identified important authors, significant works, and active institutions. As social media continues to shape advertising and user behavior, our study helps us navigate the path of SMA analysis and its importance for both academia and industry.

CONCLUSION

In conclusion, this bibliometric analysis illuminates the dynamic and rapidly evolving landscape of social media advertising. A comprehensive review of scholarly literature provided insights into various facets of this phenomenon, spanning its technical foundations to its profound impact on consumer behavior, brand engagement, and strong interconnectedness. As the field expands, it becomes evident that integrating different disciplines, methodologies, and strategies will enrich our understanding of the complex role social media advertising plays in shaping modern marketing strategies and its implications for businesses and consumers. This bibliographic analysis not only contributes to existing knowledge but also lays the foundation for future insights that will undoubtedly steer this vibrant and inevitable field into uncharted territories.

Practical Implications and Future Recommendations

More implications and future recommendations emerge from our comprehensive analysis of social media advertising (SMA). First, increasing research activity and collaboration suggests an increasing recognition of the importance of SMA. However, its relative newness in academia means there are many opportunities for further research. The geographical classification highlights the global reach of SMA research and allows for the development of different perspectives. To move forward, scholars should delve into the unexplored context of India, bridging the research gap. Examining the evolving impact of emerging technologies and approaches to data management in SMA





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can provide valuable insights into industry practices. Additionally, examining the interactions between social media platforms, advertising effectiveness, and user behavior change will enhance our understanding of this dynamic landscape Education and industry network collaboration can give us practical advantages, steering businesses towards better use of social media advertising. As the SMA field evolves in an ever-changing digital landscape, these implications and future recommendations guide researchers and practitioners toward informed and impactful contributions. The research's conclusions can help researchers and social media marketers improve their strategy creation. Researchers and SM marketers can improve strategy design by using the research's conclusions. SM marketers can better understand the market and their target audience by implementing social media analysis.

Limitations

Despite, the investigation being thorough and rigorous, the study utilizes only the Scopus database for extracting relevant information whereas the Web of Science could have been also included in collecting research literature. Secondly, while the database is being filtered before export, some essential research can be overlooked. Thirdly, although to conduct a methodical, theory-driven study on SMA analysis is applied, however, quantitative text mining can also be used.

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Table 1: Search String Used for the Study

ELEMENT	DESCRIPTION
TITLE-ABS-KEY ("social media advertising")	Search for the phrase in the title, abstract,
TITLE-AD3-RET (Social media advertising)	or keywords.
(LIMIT-TO (SUBJAREA, "BUSI") OR LIMIT-TO (SUBJAREA, "SOCI")	Restrict to specific subject areas: Business,
OR LIMIT-TO (SUBJAREA, "PSYC") OR LIMIT-TO (SUBJAREA,	Sociology, Psychology, Arts, and
"ARTS") OR LIMIT-TO (SUBJAREA, "MULT"))	Multidisciplinary.
(LIMIT-TO (DOCTYPE, "ar"))	Limit the search to articles.
(LIMIT-TO (SRCTYPE, "j"))	Restrict results to journal articles.
(LIMIT-TO (LANGUAGE, "English"))	Narrow the search to papers in English.
(LIMIT-TO (Yr>2014 to <2023))	Limit the years from 2014 to 2023

Table 2:Trends in Publication

YEAR	ARTICLES	YEAR	ARTICLES
2014	79	2019	164
2015	82	2020	161
2016	84	2021	179
2017	92	2022	198
2018	116	2023	117

Table 3: Most Productive Countries in the Domain

RANK	COUNTRY	DOCUMENTS	CITATIONS	CONTINENT	DEVELOPING/ DEVELOPED
1	UNITED STATES	90	1740	NORTH AMERICA	DEVELOPED
2	INDIA	28	303	ASIA	DEVELOPING
3	CHINA	20	147	ASIA	DEVELOPING
4	AUSTRALIA	16	212	OCEANIA	DEVELOPED
5	SOUTH KOREA	16	769	ASIA	DEVELOPED
6	UNITED KINGDOM	16	254	EUROPE	DEVELOPED
7	MALAYSIA	14	144	ASIA	DEVELOPED
8	NETHERLANDS	11	571	EUROPE	DEVELOPED
9	CANADA	9	63	NORTH AMERICA	DEVELOPED
10	GERMANY	9	116	EUROPE	DEVELOPED

Table 4: Country Collaboration Analysis

FROM	TO	FREQUENCY
USA	KOREA	8
USA	CHINA	7
GERMANY	NETHERLANDS	3
KOREA	NEW ZEALAND	3
USA	AUSTRALIA	3
USA	GERMANY	3
USA	NEW ZEALAND	3
USA	UNITED KINGDOM	3
AUSTRALIA	CANADA	2
AUSTRALIA	GHANA	2





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Table 5:Corresponding Author's Countries

RANK	COUNTRY	ARTICLES	SCP	MCP
1	USA	57	47	10
2	CHINA	21	10	11
3	INDIA	16	16	0
4	KOREA	11	5	6
5	AUSTRALIA	9	6	3
6	MALAYSIA	7	5	2
7	UNITED KINGDOM	6	3	3
8	GERMANY	5	3	2
9	HONG KONG	5	4	1

Table 6: The Most Productive Authors in SMA Analysis

RANK	AUTHOR	DOCUMENTS	CITATIONS
1	RANA M, ARORA N.	3	15
2	ARORA T, AGARWAL B.	2	50
3	BOATENG H, OKOE A.F.	2	120
4	NATARAJAN T, BALAKRISHNAN J, BALASUBRAMANIAN S.A, MANICKAVASAGAM J.	2	26
5	ADAMOPOULOS P, GHOSE A, TODRI V.	1	92
6	ADETUNJI R.R, RASHID S.M, ISHAK M.S.	1	18
7	"AHMAD A.H, IDRIS I, MASON C, CHOW S.K."	1	14
8	ALALWAN A.A.	1	278
9	ALI Y., SHAH Z.A., KHAN A.U.	1	13
10	ARORA T., AGARWAL B., KUMAR A.	1	26

Table 7: Top Authors According to Citations Per Publication

RANK	AUTHOR	DOCUMENTS	CITATIONS
1	"LEE D., HOSANAGAR K., NAIR H.S."	1	399
2	"VOORVELD H.A.M., VAN NOORT G., MUNTINGA D.G., BRONNER F."	1	321
3	LEE J., HONG I.B.	1	302
4	ALALWAN A.A.	1	278
5	ZHU YQ., CHEN HG.	1	206
6	HWANG K., ZHANG Q.	1	195
7	DUFFETT R.G.	1	177
8	JUNG AR.	1	134
9	BOATENG H., OKOE A.F.	2	120
10	ZHANG J., MAO E.	1	116





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Table 8:Co-Citations Analysis of Top 10 Authors

RANK	AUTHOR	CITATIONS	TOTAL LINK STRENGTH
1	HAIR J.F.	279	7736
2	LAW R.	252	4173
3	SARSTEDT M.	230	6622
4	RINGLE C.M.	222	6523
5	WANG Y.	196	3984
6	AJZEN I.	192	4929
7	FORNELL C.	187	4984
8	LEE J.	186	4408
9	KOTLER P.	185	2601
10	LI H.	170	3738

Table 9: Most Cited Publications

RANK	AUTHORS	YEAR	TITLE	SOURCE	CITED BY
1	LEE D.; HOSANAGAR K.; NAIR H.S. (2018)	2018	ADVERTISING CONTENT AND SOCIAL MEDIA: EVIDENCE FROM FACEBOOK	MANAGEMENT OF SCIENCE	399
2	VOORVELD H.A.M.; VAN NOORT G.; MUNTINGA D.G.; BRONNER F. (2018)	2018	ENGAGEMENT WITH SOCIAL MEDIA AND SOCIAL MEDIA ADVERTISING: THE DIFFERENTIATING ROLE OF PLATFORM TYPE	JOURNAL OF ADVERTISING	321
3	LEE J.; HONG I.B. (2016)	2016	PREDICTING POSITIVE USER RESPONSE TO SOCIAL MEDIA ADVERTISING: THE ROLE OF EMOTIONAL APPEAL INFORMATIVENESS, AND CREATIVITY	INTERNATIONAL JOURNAL OF INFORMATION MANAGEMENT	302
4	ALALWAN A.A. (2018)	2018	INVESTIGATING THE IMPACT OF SOCIAL MEDIA ADVERTISING FEATURES ON CUSTOMER PURCHASE INTENTION	INTERNATIONAL JOURNAL OF INFORMATION MANAGEMENT	278
5	ZHU YQ.; CHEN HG. (2015)	2015	SOCIAL MEDIA AND HUMAN NEED SATISFACTION: IMPLICATIONS FOR SOCIAL MEDIA MARKETING	BUSINESS HORIZONS	206

Table 10:Leading Journals in Domain

RANK	SOURCE	DOCUMENTS	CITATIONS	AVERAGE CITATION PER PUBLICATION
1	JOURNAL OF RESEARCH IN INTERACTIVE MARKETING	16	461	28.8125
2	INTERNATIONAL JOURNAL OF INTERNET MARKETING AND ADVERTISING	9	28	3.111111





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3	JOURNAL OF INTERACTIVE ADVERTISING	8	81	10.125
4	COMPUTERS IN HUMAN BEHAVIOR	6	531	88.5
5	FRONTIERS IN PSYCHOLOGY	6	30	5
6	INTERNATIONAL JOURNAL OF ADVERTISING	6	149	24.83333
7	INTERNET RESEARCH	6	353	58.83333
8	JOURNAL OF BUSINESS RESEARCH	6	81	13.5
9	JOURNAL OF RETAILING AND CONSUMER SERVICES	6	52	8.666667
10	JOURNAL OF DIGITAL AND SOCIAL MEDIA MARKETING	5	5	1

Table 11:Annual Production of Leading Journals

Year	JOURNAL OF RESEARCH IN INTERACTIVE MARKETING	INTERNATIONAL JOURNAL OF INTERNET MARKETING AND ADVERTISING	JOURNAL OF INTERACTIVE ADVERTISING	COMPUTERS IN HUMAN BEHAVIOR	FRONTIERS IN PSYCHOLOGY
2014	0	1	0	0	0
2015	1	1	0	0	0
2016	2	1	0	1	1
2017	3	1	0	2	2
2018	4	1	0	3	2
2019	6	1	1	4	2
2020	8	1	2	4	2
2021	11	5	4	5	3
2022	14	9	7	6	5
2023	16	9	8	6	6

Table 12: The Most Productive Institutions

RANK	ORGANIZATION	DOCUMENTS	CITATIONS
1	DEPARTMENT OF MARKETING, CAPE PENINSULA UNIVERSITY	2	177
1	OF TECHNOLOGY, CAPE TOWN, SOUTH AFRICA	2	1//
2	DEPARTMENT OF MARKETING, UNIVERSITY OF PROFESSIONAL	2	173
۷	STUDIES, ACCRA, GHANA	2	1/3
	"DEPARTMENT OF RETAIL, HOSPITALITY AND TOURISM		
3	MANAGEMENT, UNIVERSITY OF TENNESSEE, KNOXVILLE, TN,	2	59
	UNITED STATES"		
4	LOYOLA UNIVERSITY CHICAGO, CHICAGO, IL, UNITED STATES	2	54
	"AL-BALQA APPLIED UNIVERSITY, AMMAN COLLEGE OF		
5	BANKING AND FINANCIAL SCIENCES, AMMAN, SALT, 19117,	1	278
	JORDAN"		

Table 13: Co-Occurrence Frequency of Keywords

RAN	١K	KEYWORD	OCCURRENCES
1		SOCIAL MEDIA	88
2		SOCIAL MEDIA ADVERTISING	72





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3	ADVERTISING	46
4	MARKETING	25
5	PURCHASE INTENTION	24
6	HUMAN	20
7	FACEBOOK	19
8	SOCIAL NETWORKING (ONLINE)	18
9	SOCIAL MEDIA MARKETING	16
10	ARTICLE	14



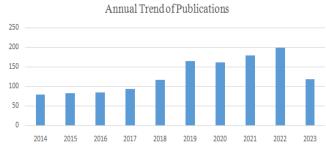
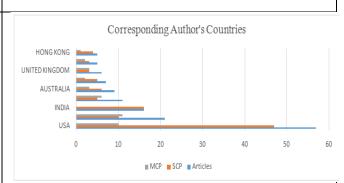


Figure 2: Annual Trend of Publications

Figure 1: Main Information About Dataset Used in Study

Country Collaboration Map



Most Prolific Sources

**DURNAL OF DIGITAL AND SOCIAL MEDIA MARKETING |

JOURNAL OF RETAILING AND CONSUMER SERVICES |

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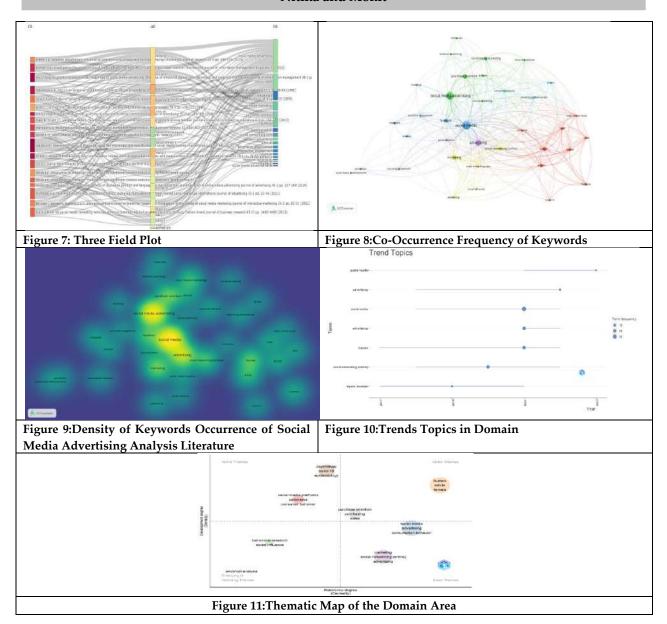
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Figure 5: Most Prolific Sources





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RESEARCH ARTICLE

A Study of Prevalence of Occupational Stress among Doctors and its Impact on Job Performance

M.Nisa¹, M.Saranyadevi², M.S.Kamalaveni³ and K.Vaitheeswari Menaga⁴

- ¹Assistant Professor, Bharathiar University PG Extension and Research Centre, Erode, Tamil Nadu, India.
- ²Research Scholar, Bharathiar University PG Extension and Research Centre, Erode Tamil Nadu, India.
- ³Associate Professor, Department of Management Studies, Sona College of Technology (Affiliated to Anna University, Chennai) Salem, Tamil Nadu, India.
- ⁴Ist MBA, Sona College of Technology, (Affiliated to Anna University, Chennai) Salem, Tamil Nadu, India.

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*Address for Correspondence

M.Nisa

Assistant Professor, **Bharathiar University** PG Extension and Research Centre, Erode, Tamil Nadu, India.



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ABSTRACT

Doctors plays a vital role in the Country especially house officers. They are under a great deal with stress variety of occupational stressors. Occupational stressors which may include to organizational inefficiency, high staff turnover, absenteeism due to sickness, decreased quality of practice, increased costs of health care and decreased job satisfaction. The most of the organizational outcomes they are affected by occupational stress and its job performance. The purpose of study was to investigate the effect of job stress on job performance Government doctors in Salem District, Tamil Nadu The universal study on Government doctors in Salem District. The targeted respondents which were present at that time were used in the 520. The data obtained through descriptive statistics, Spearman's Correlation and Multi Regression. The analysis showed strong support for the hypothesis that there is an inverse relationship between job stress and job performance indicating that there is high job stress in the house officers, resulting in low job performance. Appropriate stress management should start from improved health and good interpersonal relationships. The study reveals that job performance is impeded by the stress at workplace. The prevention and management of workplace stress requires both individual and organizational level interventions, as it a net result of factors causing stress and the individual's adaptation to the stressors.

Keywords: Occupational stressors; workplace stress, job performance and workplace stress





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INTRODUCTION

The current turbulent environment in the health care field requires doctors and 150 organizations to reexamine their practices. Medicine is an inherently stressful profession with long working hours, ethical dilemmas, difficult patients and conflicting demands. Professionally, in true sense, the doctors are on 24-hour duty. Many physicians and surgeons work long, irregular hours; over one-third of full-time physicians worked 60 or more hours a week in 2023. The physical and; psychological demands of the profession often make physicians more vulnerable to high levels of stress. The effects of stress job practice are evidenced as increased errors in prescribing, limited team working, more patients' complaints and sickness absence. Stress has been defined in different ways over the years. Originally, it was conceived as pressure from the environment, then as strain within the person. The generally accepted definition today is one of the interactions between the situation and the individual. It is the psychological and physical are not sufficient to cope with demands and pressures of the situation. Thus, stress is more likely in some situations than others and in some individuals than others. Stress is not always negative or harmful and indeed, the absence of stress is death.

Stress is the non-specific response of the body to any demand; health sector is one of the most stressful professions and pointed out the necessity of considering and investigating occupational stress, since performance declines under stressful situations. Therefore, stress at workplace becomes a concern to organization administrators. Several authors attributed the lack of progress in the areas of stress research 151 organizations to the fact that stress seemed to be related to such a large number of conditions which prevented a systematic focus. Beehr used a very general definition in which "anything about an organizational role that produces adverse consequences for the individual" was called role stress. They proceeded to the conclusion that a conditions termed role overload was viable and this correlated positively with job stress. Stress indicators to role ambiguity in the study indicated low motivation to work. This study was based on a sample of 651 persons, including 213 from service departments of a hospital. The primary source of source cited by respondents indicated juggling multiple roles, having young children, time issues (too much work, too little time changing practices patterns. Job performance is the result of three factors working together: skill, effort and the nature of work conditions. Skills include knowledge, abilities and competencies the employee brings to the job, effort is the degree of motivation the employee puts forth toward getting the job done; and the nature of work conditions is the degree of accommodation of these conditions in facilitating the employee's productivity.

CONCEPTUAL MODEL

A conceptual framework speaks to the researcher's union of writing on the most proficient method to clarify a marvel. It maps out the activities required over the span of the investigation given his past learning of other researchers' perspective and his perceptions regarding the matter of research. At the end of the day, the conceptual framework is the researcher comprehends of how the specific factors in his examination associate with each other. In this manner, it recognizes the factors required in the examination. It is the researcher's "guide" in seeking after the examination.

SCALES USED

All items are measured on 5 –point Likert scale (5-strongly agree to 1 strongly disagree). Few items are measured on 6 point and 4 point scale.

PILOT TESTA

The pilot study was conducted among 520 members of Government doctors in Salem District i.e. 10 per cent of the total sample to check the feasibility and reliability of the research schedule. In the light of the experience gained in the pilot study the research schedule has been modified to suit the sample groups and finalized to conduct the survey. In the light of the experience gained in the pilot study the research schedule has been modified to suit the sample groups and finalized to conduct the survey. The perception of the respondents has been tested for its





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reliability using Cronbach's Alpha. The value obtained was 0.921, which shows that instrument is highly reliable. The distribution curve is normal. Based on the pilot study necessary modifications were made in the interview schedule. The researcher sought adequate guidance through the research supervisor.

SAMPLING DESIGN

According to *Politand Hungler* (1999), "Sampling refers to the process of selecting a portion of the population to represent the entire population". The representative sample consists of subsets of the elements of a population which allows for the study results to be generalized. The characteristics of the sample population are intended to be representative of the target population. This study incorporates stratified random sampling, as the population of Government doctors in Salem District were taken up for the study. The entire population of Government doctors in Salem District was classified into four strata and from each stratum samples were drawn through simple random sampling method.

POPULATION AND THE STUDY AREA

The sample population taken from the Government doctors in Salem District with regards to Government doctors and the study area is occupational stress among the Government doctors and how the stress is affecting the individual doctor serving in different categories and their reaction towards the stress that they undergone during their services. This study area incorporates in Salem District and focuses on doctor's occupational stress and its impact on the job performance and job satisfaction.

SAMPLING TECHNIQUES

Stratified random sampling techniques are deployed in this study. The entire geographic area of Salem District has been divided into four parts or strata or group of police stations. From each strata or group or part samples were drawn on simple random and totally 520 Government doctors were selected through the stratified random sampling as equal proportion of samples were selected from each strata.

SAMPLE SIZE

This study is carried out with the sample size of 520 across the various categories of Government doctors.

DATA COLLECTION

Data collection was carried out with the help of questionnaire from the employees of Government doctors primary data collection was carried out with the help of questionnaires Government doctors. The field survey technique was employed to collect the pertinent data from 520 Government doctors in the study area. Interview schedule was the main tool for collecting the primary data. Much effort was taken to prepare the interview schedule in a systematic way by designing adequate and relevant questions to ensure better achievement of the research objective. The interaction technique is applied to collect certain relevant data in order to facilitate the study. The Primary data was supplemented by spate of secondary data. The secondary data pertaining to the study was gathered from the various sources pertaining to Government doctors their stress experienced by doctors. Further, the secondary data were collected from leading journals and a number of standard reference books were referred to obtain pertinent literature on occupational stress of Government doctors.

DATA ANALYSIS FRAMEWORK

After analyzing the various data collection methods and research instruments, a questionnaire having questions with multiple-choice responses and a 5 point 'Likert-type scale' with -2 being Strongly Disagree and2 being Strongly Agree, was selected as the survey instrument. The present study is descriptive cum exploratory in nature. The study is descriptive as it describes the occupational stress of Government doctors in Salem District. After analyzing the various data collection methods and research instruments, an questionnaire having questions with multiple-choice responses and a 5 point 'Likert-type scale' with 1 being strongly disagree and 5 being Strongly Agree, was selected as the survey instrument. Data entry, processing and analysis were done using SPSS for Windows (Version 22.0)





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spreadsheet program and Microsoft Excel 2007. Descriptive statistics (frequencies, scores, mean, maximum, minimum) were determined. The actual processing and analysis started with data cleaning to remove the gaps and ensure consistency. In order to test the association between independent variables and dependent variables, chi-square test was applied; Correlation, Factor Analysis, ANOVA,'t' test model was applied to find out the variation within samples and between samples. The present study is descriptive cum exploratory in nature. The study is descriptive as it describes the occupational stress of Government doctors in Salem District. It also seeks to explore and measure the level of stress the Government doctors.

SAMPLING FRAME

The sample was designed in a two phases. In the first phase the Government doctors in Salem District was selected. In second phase the respondents were selected from Government doctors in Salem District. The list of the police stations covered in the district, which served as the sampling frame for the purpose of the study. This provided the complete list and various Government doctors serving at Government doctors were taken into consideration from the same. The above table mentioned table 1 shows the frequency distribution of based on the socio economic characteristics of the Government doctors in Salem District. The highest Majority of the respondents are male (79.4%) of the Government doctors. 20.8% percent of respondents fall in the age group of 31-35 years, 41.2% of the respondents educational qualification is professional, In respect of the marital status majority of the respondents are married persons (25%), based on the Number of dependents of the respondents the 5 to 6 category is 40%, 36.3% of the respondents are living in rural area. Majority of the respondents designation is head constable 26.9%, Based on the Monthly Income of the respondents the 31,000-60,000 32.9% of Government doctors, In respect of the Number of years of experience as the Government doctors 33.3%, percent respondents falls under the category of 3-5 years as experience in police department. Based on the transferred of the respondents 1-5 times 31.7% of Government doctors 33.5% percentage of Government doctors undergone training programs in 6-10 times. From the above table, the highest mean value meant for Lack of training on new equipment is the most important factor as considered by the Government doctors that led to severe occupational stress followed by Finding time to stay in good physical condition is next important factor that led to occupational stress amongst the Government doctors in Salem District. The Government doctors perception that Risk of being injured on the job was the next important factor that led to occupational stress followed by Fatigue (shift work and over time etc.,) and Staff shortages were considered as the occupational stress of the Government doctors in Salem District.

The other factors responsible the occupational stress were Shift duty work, Working alone at night, Over time demands, Risk of being injured on the job, Work related activities on days off (e.g. law and order, community events), Traumatic events (e.g. MVA, domestic, death, injury), Managing social life outside of work, Not enough time available to spend with friends and family, Enormous paper work, Poor cope up with work, Dealing with coworkers, The feeling that different rules apply to different people (e.g. favoritism) Feeling like the police person always have to prove himself or herself to the organization, Excessive administrative duties, Constant changes in policy / legislation, Bureaucratic red tape, Too much computer work, Perceived pressure to volunteer free time, Limitations to social life and the Negative comments from the public were responsible for occupational stress of the Government doctors in Salem District There is significant relationship between the mean score of age of the Government doctors and associated risk factor of occupational stress of Government doctors. Shift duty work Managing social life outside of work, Poor cope up with work, Feeling like you always have to prove yourself to the organization, Constant changes in policy / legislation, Staff shortages, Bureaucratic red tape, Lack of training on new equipment, Too much computer work, Limitations to your social life (eg. Who your friends are, where you socialize) and Fatigue (shift work and over time etc.,) were the factors causing occupational stress and significant at 1% level. Therefore it is concluded that there is significant relationship between the mean score of age of the Government doctors and associated risk factor of occupational stress of Government doctors. Work related activities on days off (e.g. law and order, community events), the feeling that different rules apply to different people and Perceived pressure to volunteer free time were significant at 5% level. The above table clearly indicated that the Shift duty work is the only factor considered by the Government doctors that cause occupational stress amongst they and it is significant at 1% level. Therefore we conclude that there is significant relationship between the mean score of





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qualification of the Government doctors and shift duty was considered as the most important risk factor of occupational stress of Government doctors. It is evident that the as the Government doctors becomes more qualified they would manage their occupational stress very well. It is evident that the higher the qualification of the Government doctors the lower is the occupational stress amongst the Government doctors. From the above table, it is found out that all the variables related to the risk factor of occupational stress of Government doctors had significance value less than 0.05 at 1 Per cent significance, thus the null hypothesis is rejected. Thus, it is concluded that there is significant difference between mean ranks towards the risk factor of occupational stress of Government doctors. Out of the twenty four risk factor of occupational stress of doctors the "Lack of training on new equipment" has the highest mean rank (14.21). Therefore this risk factor associated with occupational stress of the Government doctors. High value of KMO (0.530> .05) of indicates that factor analysis is useful for the present data. The significant value for Bartlett's test of Sphericity is 0.000 and is less than 0.05 which indicates that there exists significant relationships among the variable. The result value of KMO test and Bartlett's test indicates that the present data is useful for factor analysis. From tree analysis model summary "Feeling like you always have to prove yourself to the organization. Finding time to stay in good physical condition is important independent variable.

From the Tree structured analysis, it is determined that out of the twenty four attributes of risk factor of occupational stress of Government doctor, the most influencing factor is identified as "Feeling like you always have to prove yourself to the organization. This may be due to the fact that, any Government doctor professional should possess some fundamental occupational stress strategies rendered to their employees. Therefore, to risk factor of occupational stress of Government doctor, the above said factor "Feeling like you always have to prove yourself to the organization. Finding time to stay in good physical condition is identified as the most important independent variable. Here the coefficient of Risk Factor is 0.107 represents the partial effect of Risk factor on Job Performance, holding the other variables as constant. The estimated positive sign implies that such effect is positive that Job performance would increase by 0.107 for every unit increase in risk factor and this coefficient value is significant at 1% level. The coefficient of organizational climate is 0.007 represents the partial effect of Organizational climate on Job Performance, holding the other variables as constant. The estimated negative sign implies that such effect is negative that Job Performance would decrease by 0.007 for every unit decrease in job performance and this coefficient value is no significant at 1% level. The coefficient of Occupational Stress is .263 represents the partial effect of occupational stress on Job Performance, holding the other variables as constant.

The estimated positive sign implies that such effect is positive that job performance would increase by .263 for every unit increase in Occupational Stress and this coefficient value is significant at 1% level. The coefficient of Stress level is 443 represents the partial effect of Stress level on job performance, holding the other variables as constant. The estimated positive sign implies that such effect is positive that job performance would increase by .443 for every unit increase in stress level and this coefficient value is significant at 1% level. The coefficient of risk factor is .038 represents the partial effect of risk factor on Prevention of the Stress, holding the other variables as constant. The estimated negative sign implies that such effect is negative that prevention of the stress would increase by .038 for every unit decrease in risk factor and this coefficient value is significant at 1% level. The coefficient of organizational climate is 0.009 represents the partial effect of organizational climate on prevention of the stress, holding the other variables as constant. The estimated negative sign implies that such effect is negative that prevention of the stress would decrease by 0.009 for every unit decrease in organizational climate and this coefficient value is significant at 1% level. The coefficient of Occupational stress is .169 represents the partial effect of occupational stress on prevention of the stress, holding the other variables as constant. The estimated positive sign implies that such effect is positive that prevention of the stress would increase by .169 for every unit increase in occupational stress and this coefficient value is significant at 1% level. The coefficient of stress level is 0.007 represents the partial effect of stress level on prevention of the stress, holding the other variables as constant. The estimated negative sign implies that such effect is negative that prevention of the stress would decrease by 0.007 for every unit decrease in stress level and this coefficient value is significant at 1% level. The coefficient of Job performance is 2.068 represents the partial effect of job performance on Job satisfaction, holding the other variables as constant. The estimated positive sign implies that such effect is positive that Job satisfaction would increase by 2.068 for every unit increase in job performance and





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this coefficient value is significant at 1% level. The coefficient of Prevention of the stress is 3.701 represents the partial effect of prevention of the stress on Job satisfaction, holding the other variables as constant. The estimated positive sign implies that such effect is positive that Job satisfaction would increase by 3.701 for every unit increase in prevention of the stress and this coefficient value is significant at 1% level. Based on standard coefficient, Prevention of the stress (1.246) is most important variable of Job satisfaction followed by Job performance (0.987) and stress level (0.359). From the above table it is found that the calculated P value is 0.015 which is less than 0.05 which indicates perfectly fit. Here GFI (Goodness of Fit Index) value and AGFI (Adjusted Goodness of Fit Index) value is greater than 0.9 which represent it is a good fit. The calculated CFI (Comparative Fit Index) value is 0.996 which means that it is a perfectly fit and also it is found that RMR (Root Mean Square Residuals) is 0.155 and RMSEA (Root Mean Square Error of Approximation) value is 0.078 which is less than 0.08 Which indicated it is perfectly fit.

IMPACT OF STRESS ON DOCTORS

Several studies have shown that occupational stress can lead to various negative consequences for the individual and the workplace (Oginska-Bulik, 2006) in healthcare sector. Stress produces a range of undesirable, expensive, and debilitating consequences (Ross, 2005), which affect both individuals and hospitals. Early individual behavioral reactions may include onset or increased smoking and alcohol use. Individuals may tend to keep late nights in clinics/ offices without accompanying increased productivity. While others might become irritable, some will tend to intense seclusion and individualism. Many studies deal with the topic area sleep loss tiredness exhaustion work stress burnout. They mostly come to the same conclusion, that stress has a major influence in doctors working lives but also their personal lives. Stress and fatigue may result in increased insecurity in clinical decisions and may therefore negatively affect the ability to practice medicine adequately, responsibly, and without error. 50% of such incidents resulted in a loss in treatment standards; 7.4 % were expressed in serious treatment errors, although actual death had been avoided.

CONCLUSION

Overall, the study reveals that majority of Government Doctors agree significantly that they experienced occupational-stress. It also supports the assumption that health sector employees are among the highest groups subjected to work stress. Results also indicated that prevalence of occupational-stress was not influenced by gender, age and experience of the employee. The stress in turn affects the performance of the individual. Thus the policy makers must pay the attention towards the issue of occupational stress in hospitals through problem recognition and problem-solving activities to deal with the sources of stress so that effective and efficient performance of the doctors can be ensured.(The stress of the Government doctors measured by analyzing the dimensions considered for the. Study with the help of the hypothetical statements and statistical tools considered for the study. The study reveals that a part of the conceptualized research model was empirically proved. These findings are interpreted in the final chapter for future research and Government doctors)

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Table 1

Demography and Socio Economic	Variables	Frequency	Percent
	Male	413	79.4
Gender	Female	107	20.66
	Total	520	100.0
	21-26	103	19.8
	27-30	107	20.6
Ago	31 –35	108	20.8
Age	35-40	103	19.8
	Above 45	99	19.0
	Total	520	100.0
Education Qualification	Up to HSC	52	10.0
Education Qualification	Diploma	91	17.5





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	Graduation	83	16.0
	P.G	80	15.4
	Professional	214	41.2
	Total	520	100.0
	Married	130	25.0
	Un Married	171	32.9
Status Marital	Divorced	121	23.3
	Widow	71	13.7
	Total	520	100.0
	Up to 2	60	11.5
	3 to 4	164	31.5
Number of Dependents	5 to 6	208	40.0
	above 7	85	16.3
	Total	520	100
	Urban	145	27.9
DI (:1	Rural	189	36.3
Place of residence	Semi urban	186	35.8
	Total	520	100.0
	Medical officer	87	16.7
	Senior Medical Officer	140	26.9
	Chief Medical Officer	133	25.6
Designation	General Duty Medical officer	91	17.5
	Inspector Medical Officer	69	13.3
	Total	520	100.0
	Up to 30,000	110	21.2
	31,000-60,000	171	32.9
Income	61,000- 90,000	144	27.7
	Above 91,000	95	18.3
	Total	520	100.0
	Below 2 year	116	22.3
	3-5 year	173	33.3
	6-15 year	138	26.5
	16-30 year	93	17.9
	Total	520	100.0
TATaula as a said a said	No	134	25.8
Work experience	1-5	165	31.7
	6 – 15	122	23.5
	Above 15	99	19.0
	Total	520	100.0
	1-5	96	18.5
	6 – 10	174	33.5





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11 – 25	162	31.2	
Above 25	88	16.9	
Total	520	100.0	

Table -2 Mean and standard deviation for risk factor of occupational stress of Government doctors

Risk factor of occupational stress of Government doctors	Mean	Std. Deviation
Shift duty work		1.210
Working alone at night	3.43	1.328
Over time demands	3.49	1.277
Risk of being injured on the job	3.64	1.284
Work related activities on days off	3.18	1.314
Traumatic events	3.43	1.290
Managing social life outside of work	3.24	1.236
Not enough time available to spend with friends and family	3.37	1.325
Enormous paper work	3.49	1.339
Poor cope up with work	3.49	1.304
Dealing with co-workers	3.54	1.322
The feeling that different rules apply to different people	3.47	1.333
Feeling like you always have to prove yourself to the organization.	3.53	1.289
Excessive administrative duties		1.324
Constant changes in policy / legislation		1.287
Staff shortages		1.291
Bureaucratic red tape	3.59	1.328
Too much computer work	3.55	1.306
Lack of training on new equipment		1.362
Perceived pressure to volunteer free time		1.283
Finding time to stay in good physical condition		1.302
Limitations to your social life		1.286
shift work and over time etc.,	3.64	1.232
Negative comments from the public	3.54	1.278

Source: Primary Data Generated from the Government doctors

Table-3 ANOVA test for age of the Government doctors and associated risk factor of occupational stress of Government doctors

Risk factor of occupational stress of Government doctors	F-Value	Sig. Value
Shift duty work	32.225	.000**
Working alone at night	.134	.970
Over time demands	2.802	.025
Risk of being injured on the job	2.827	.024
Work related activities on days off		.010*
Traumatic events (e.g. MVA, domestic, death, injury)		.041
Managing social life outside of work		.000**
Not enough time available to spend with friends and family		.024
Enormous paper work 2.174 .0		





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Poor cope up with work	3.805	.005**
Dealing with co-workers	1.188	.315
The feeling that different rules apply to different people	3.149	.014*
Feeling like you always have to prove yourself to the organisation.	9.377	.000**
Excessive administrative duties	2.446	.046
Constant changes in policy / legislation	3.838	.004**
Staff shortages	3.671	.001**
Bureaucratic red tape		.005**
Too much computer work		.000**
Lack of training on new equipment		.000**
Perceived pressure to volunteer free time		.024*
Finding time to stay in good physical condition		.057
Limitations to your social life		.003**
Fatigue (shift work and over time etc.,)		.000**
Negative comments from the public	1.678	.154

Source: Primary Data Generated from the employees of Police department Note : * Denote at 5% level - **Denote at 1% level

Table -4 ANOVA test for qualification of the doctors risk factors of occupational stress of Government doctors

Risk factors of occupational stress of Government doctors	F-Value	Sig.Value
Shift duty work	9.432	.000**
Working alone at night	1.819	.124
Over time demands	.601	.662
Risk of being injured on the job	.470	.758
Work related activities on days off	.218	.929
Traumatic events (e.g. MVA ,domestic , death , injury)	.518	.722
Managing social life outside of work	.552	.698
Not enough time available to spend with friends and family	1.096	.358
Enormous paper work	1.067	.372
Poor cope up with work		.016
Dealing with co-workers		.019
The feeling that different rules apply to different people		.122
Feeling like you always have to prove yourself to the organization		.406
Excessive administrative duties	1.592	.175
Constant changes in policy / legislation	.620	.649
Staff shortages		.289
Bureaucratic red tape	2.088	.081
Too much computer work		.027
Lack of training on new equipment	.477	.753
Perceived pressure to volunteer free time	.631	





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Finding time to stay in good physical condition		.010
Limitations to your social life		.719
Fatigue		.808
Negative comments from the public		.048

Source: Primary Data Generated from the employees of Police department Note: * Denote at 5% level - **Denote at 1% level

Table -5 Friedman test for significant difference between mean ranks towards the risk factor of occupational stress of Government doctors

risk factor	Mean Rank	Chi Squar e	DF	Significant
Shift duty work	12.21			
Working alone at night	12.00			
Over time demands	12.47			
Risk of being injured on the job	13.23			
Work related activities on days off	10.84			
Traumatic events	12.10			
Managing social life outside of work	11.25			
Not enough time available to spend with friends	11.82			
Enormous paper work	12.56			
Poor cope up with work	12.46			
Dealing with co-workers	12.72			
The feeling that different rules apply	12.26			
Feeling like you always have to prove your self	12.62			
Excessive administrative duties	12.84			
Constant changes in policy / legislation	11.79			
Staff shortages	13.10			
Bureaucratic red tape	12.90	130.582	23	0.000**
Too much computer work	12.66			
Lack of training on new equipment	14.21			
Perceived pressure to volunteer free time	12.68			
Finding time to stay in good physical condition	13.32			
Limitations to your social life	12.11			
Fatigue (shift work and over time etc.,)	13.22			





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Negative comments from the public	12.64		

Source: Primary Data Generated from the employees of Government doctors Note: * Denote at 5% level - **Denote at 1% level

Table -6 KMO and Bartlett's Test of risk factor of occupational stress of Government doctor

KMO and Bartlett's Test			
Kaiser-Meyer-Olkin Measure of Sampling Adequacy .530			
	Approx. Chi-Square	1613.290	
Bartlett's Test of Sphericity	Degree of freedom	276	
1 ,	Significance	.000**	

Source: Primary Data Generated from the employees of occupational stress of doctors. Note: **Denote at 1% level

Table – 7 Model summary for risk factor of occupational stress of Government doctor

	Growing Method	CHAID
	Dependent Variable	Overall Job satisfaction of the Government doctors
Specifications	Independent Variables	Shift duty work, Working alone at night, Over time demands, Risk obeing injured on the job, Work related activities on days off (e.g. law and order, community events), Traumatic events (e.g. MVA, domestic, death, injury), Managing social life outside of work, Not enough time available to spend with friends and family, Enormous paper work, Poor cope up with work, Dealing with co-workers, The feeling that different rules apply to different people (e.g. favouritism), Feeling like you always have to prove yourself to the organisation., Excessive administrative duties Constant changes in policy legislation, Staff shortages, Bureaucratic rectape, Too much computer work, Lack of training on new equipment Perceived pressure to volunteer free time, Finding time to stay in good physical condition, Limitations to your social life (eg. Who your friendare, where you socialize), Fatigue (shift work and over time etc.,) Negative comments from the public
	Validation	None
	Maximum Tree Depth	3
	Minimum Cases in Parent Node	100
Result	Minimum Cases in Child Node	50





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Inde	ependent Variables Included	Feeling like you always have to prove yourself to the organisation., Finding time to stay in good physical condition
N	Number of Nodes	7
	Number of Terminal Nodes	5
	Depth	2
N	Number of Nodes	7

Source: Primary Data Generated from the employees of Government doctor

Table – 8 Risk model for risk factor of occupational stress of Government doctor

Estimate	Std.
Estimate	Error
.039	.005
Growing Meth Dependent Variable: Overall Job satis:	

Source: Primary Data Generated from the employees of Government doctor

Table -9 Gain summary nodes for risk factor of occupational stress of Government doctor

Gain Summary for Nodes					
Node	N	Percent	Mean		
4	178	34.2%	5.00		
3	76	14.6%	4.00		
2	136	26.2%	3.00		
6	59	11.3%	1.92		
5	71	13.7%	1.66		
	•				

Growing Method: CHAID
Dependent Variable: Overall Job satisfaction of the Government doctors

Source: Primary Data Generated from the employees of Government doctor

Table -10 Number of variables in the SEM

Number of variables in your model		
Number of observed variables	7	
Number of unobserved variables		
Number of exogenous variables		
Number of endogenous variables		

Source: Output generated from AMOS 20

Table – 11 Variables in the Structural Equation Model Analysis

<u>,</u>	Unstandardize dcoefficient	S.E	Standardize d coefficient	t value	P value
Job Performance < Risk Factor	.107	.027	.138	3.924	<0.001**
Job Performance < Organizational Climate	007	.049	008	146	.884





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		,			
Job Performance < Occupational Stress	.263	.056	.273	4.712	<0.001**
Job Performance < Stress Level	.443	.048	.359	9.296	<0.001**
Prevention Of The Stress < RiskFactor	038	.017	069	-2.240	.025*
Prevention Of The Stress < Organizational Climate	009	.018	014	521	.602
Prevention Of The Stress <occupational stress<="" td=""><td>.169</td><td>.035</td><td>.248</td><td>4.859</td><td><0.001**</td></occupational>	.169	.035	.248	4.859	<0.001**
Prevention Of The Stress < Stress Level	007	.034	008	190	.849
Job Satisfaction < Job performance	2.068	.325	.987	6.353	<0.001**
Job Satisfaction < Prevention Of The Stress	3.701	1.077	1.246	3.435	<0.001**

Note: ** denotes significant at 1% level Source: Output generated from AMOS 20 $\,$

Table – 12 Model fit summary of Structural Equation Model

Indices	Value	Suggested value
Chi-square value	8.342	-
P value	0.015	> 0.05 (Hair et al.,1998)
GFI	0.995	> 0.90 (Hu and Bentler,1999)
AGFI	0.937	> 0.90 (Hair et al., 2006)
CFI	0.996	> 0.90 (Daire et al., 2008)
RMR	0.155	>0.08 (Hair et al., 2006)
RMSEA	0.078	<0.08 (Hair et al., 2006)

Source: Output generated from AMOS 20





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RESEARCH ARTICLE

Analyzing Business Crimes by Implementing OSINT Architecture

Yuthvek .M J¹, Adheena Shibu¹, Charithra Y¹ and K. Kalaiselvi^{2*}

¹Student, Department of Forensic Science, Kristujayanti College (Autonomous), (Affiliated to Bengaluru North University), Bangalore, Karnataka, India.

²Associate Professor, Department of Computer Science, Kristujayanti College (Autonomous), (Affiliated to Bengaluru North University) Bangalore, Karnataka, India.

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*Address for Correspondence K. Kalaiselvi

Associate Professor, Department of Computer Science, Kristujayanti College (Autonomous), (Affiliated to Bengaluru North University) Bangalore, Karnataka, India. Email: kalaiselvi@kristujayanti.com



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ABSTRACT

Financial losses and damage to a company are caused by business crimes. In this study, we used opensource tools, which are freely available tools, to analyze those business crimes. We started to analyze three made-up cases using 10 different OSINT tools to gain a deeper understanding of the concept. Our study's made-up cases were based on business crimes such as money laundering, insider trading, and data breaches. We have examined the effectiveness of those tools. We have obtained permission from the Te2 technique company to use them as a model. OSINT tools are very useful for gathering publicly available data, but their results are not reliable according to our study's conclusion. Only 4 out of the 10 chosen tools were helpful for analyzing those cases. The data we get from those tools tends to have more errors, so we need to be careful when using them.

Keywords: Business Crimes, Open-Source Intelligence, Digital Forensics, White-collar Crimes, OSINT Tools.

INTRODUCTION

Complex financial offences that can be committed by individuals, companies, or employees are called white-collar crimes or business crimes. The majority of crimes committed in large corporations with extensive technology and global networks involve deceit, concealment of information, or breach of trust. Because of this, these crimes have become an important international issue. Fake deals, stealing, money laundering, theft of intellectual property,





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insider trading, cybercrime, and other deceitful business practices are all part of business crime. These crimes can result in significant financial losses, damage a company's reputation, and cause legal consequences[1]. Detecting and preventing these crimes is crucial to maintain fairness in the business world and financial markets. Business crimes often involve using technology incorrectly and taking advantage of digital systems. Cyber-attacking and hacking are common types of business crime. Criminals break into digital systems in these circumstances to steal important information or disrupt business activities[2]. Corporate spying is a significant area where businesses secretly take action to advance their competitors, often by stealing ideas. The use of digital forensics is essential for detecting cases of individuals trading secrets, hiding illegal money, or cheating with money in businesses. It can be helpful in investigating when data is stolen, people's privacy is violated, or when online scams and tricks occur, which can have negative effects on businesses and individuals[3]. Digital forensics experts deal with other types of business crimes, such as stealing ideas, being corrupt, offering bribes, not following rules, and breaking regulations. The focus of this research is to investigate the methods and tools utilized in digital forensics to understand and reduce the impact of various types of business crimes. In the realm of research and investigation, there exists a technique called open-source intelligence (OSINT), or OSINT for short. By gathering information from publicly available sources, this method can be used for a variety of purposes, including learning about businesses and apprehending individuals involved in illicit activities[4]. The website of OSINT framework can be accessed via the link: https://osintframework.com/. The focus is on finding relevant information that pertains to the intelligence question and offering practical intelligence in support of an investigation. Our research encompasses a comprehensive examination of how OSINT tools work when it comes to analyzing business crimes. Our focus is on finding effective ways to use tools during investigations while utilizing open-source resources for thorough examination and analysis. In Fig 1, you can see the homepage of the OSINT framework website.

REVIEW OF LITERATURE

Cyber security can also be increased by the usage of OSINT in crimes like phishing detection, threat intelligence, hate speech detection, fake news detection, human trafficking, child trafficking, criminal profiling, etc. In this paper they have discussed the state of OSINT nowadays and its benefits, importance in cybercrime investigations. The significance of this study lies in its ability to enhance investigators' techniques and applications of OSINT[1]. The threats and vulnerabilities were calculated in this study based on the impacts of each review of the databases. The underlying research challenges that impact source intelligence are also highlighted in the paper. For this article, they have searched many online databases and reviewed 18 research papers that investigated the effects of social media technologies under the concept of OSINT [5]. In this article they have proposed a new way to handle online child abuse investigations using a theoretical model after a brief explanation of crowd sourcing and OSINT techniques. The theoretical model comprises organizational, technical and legal aspects of selected skilled and tightly grouped volunteers to help in online child abuse cases using OSINT and forensic examinations without compromising the legal procedures[6]. In this study they emphasize providing a helping hand in digital investigations with efficient tools. Here, the method they used was to analyze the activities done, the people linked, and the persons contacted for communications by a suspected employee. This study highlights how to handle digital evidence and retrieve multimedia information for analysis using OSINT[7]. Digital forensic Intelligence and Open-source Intelligence are balanced in this paper. The study analyzes the application of this framework to investigate the additional information contained in the digital forensic data subset. This highlights the prompt extraction, analysis, and automated entity identification of real-world data by Open-Source Intelligence [8]. In this study they deal with the current methods used by investigators for cybercrimes, the challenges they face, and future directions for investigations. This study highlights the use of open-source intelligence and the use of many such tools for investigation purposes. They emphasize the selection of tools for different types of cases based on the information that can be provided by each tool and its effectiveness. The study concludes that for better effective investigations, many tools must be combined and used since there is no one single tool for all the work to be done[9]. The current situations and security trends of OSINT are explained in this paper as data collected using OSINT can be used for





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malicious activities. The paper discusses the types of cybercrimes and security threats that can arise when OSINT is exploited by malicious users. Furthermore, the paper provides countermeasures, such as the proposed need for security requirements, that can be employed in the OSINT environment [10]. In this study they deal with topic modelling processes, which play a pivotal role in providing significant results that can be useful in OSINT. The study focuses on LDA (Latent Dirichlet Allocation), its processes, and its advantages, and offers suggestions for future assistance in cybercrime investigations[11]. Open-source research techniques, analytical tools, and datasets are utilized in the study to investigate organized crimes like human trafficking. The study indicates that large-scale data analytics are required to deal with more reports of cyber-facilitated crime[12]. The study relies on a comprehensive review of OSINT-related literature, its applications in cybercrime, and relevant case studies. The study implies the challenges and limitations faced by the OSINT framework and proposes potential solutions like integration of machine learning, new algorithms based on artificial intelligence and proper handling of data collection and analysis[13].

METHODOLOGY

The Process of Collecting Digital Evidence with OSINT Techniques:

- **Information Gathering:** Begin by defining the scope of the investigation and the specific information needed, such as details about a person, organization, event, or online activity.
- Tool Selection: Select the appropriate OSINT tools and resources to access and retrieve information from the specified sources. There were a variety of tools available, including search engines and specialized OSINT software and platforms.
- **Data Collection:** Use OSINT tools to gather data from various sources, such as online searches, social media, and public databases in an ethical and legal manner.
- **Data Verification:** Verify the accuracy and reliability of collected information. Verify facts and eliminate false or misleading data by cross-referencing data from multiple sources.
- Data Preservation: Safeguard digital evidence by preserving original URLs, taking screenshots, and archiving web pages as they appeared at the time of collection.
- Data Analysis: Analyze the collected data to extract insights, patterns, and connections using data manipulation, keyword searches, and visualizations.
- Documentation: It is crucial to document the entire process of collecting OSINT, including the sources utilized, search queries, and any modifications made to the data. Legal and investigative purposes require accurate documentation.

The OSINT framework and ten different tools will be used to analyze three hypothetical cases of different business crimes in this paper. We will evaluate the effectiveness of the tools and interpret the results obtained. In the end, we will present a detailed framework for investigating such cases using OSINT tools. It is important to note that all the cases mentioned in this paper are hypothetical, but for the sake of understanding, we will use Te2 Technique, a service-based company located in Tamil Nadu, India, as an example. The steps to collect digital evidence using OSINT are shown in Fig 2

CASE-1 (Money laundering)

Te2 Technique company is suspected of laundering money. Te2 Technique Corporation is a company that spans multiple countries. Manufacturing, import/export, and retail are legitimate businesses that the company is involved in. The management suspects a complicated money laundering scheme to hide the origins of illegal funds. Mr. Yuthvek, the CEO of Te2 Technique Corporation, is suspected of being the mastermind behind the money laundering operation. Mrs. Josephine, who is also the Chief Financial Officer (CFO) of the company, is being investigated because they both have the ability to oversee financial transactions.





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Questions

- 1. What is the purpose of the investigation?
- 2. Do the suspects have any suspicious communications?
- 3. Are any of their devices experiencing unusual bank transfers?
- 4. Are there any discrepancies between the financial record and the business operations that occurred?

Investigative Tools

Hunter is an OSINT tool to obtain the email addresses of companies and their employees associated with a domain. Here, it was used to determine if the email addresses of the suspects were authentic, which was then provided to the respective banks for the bank statements. The website can be accessed by using the link https://hunter.io/search.The free digital forensic software may have limited effectiveness and capabilities compared to premium options, lacking advanced features and user-friendliness, which can be a disadvantage for those who require more specialized functionality in their forensic work. The homepage of hunter tool is shown in fig 3,4. Whatismyipaddress is an OSINT tool that identifies potentially suspicious emails originating from the sender. This OSINT utility is used to analyze IP addresses associated with emails receiving money to determine hostnames and geographical location. The tool's website can be accessed through the link https://whatismyipaddress.com.The main focus of "Whatismyipaddress" is to provide information about IP addresses from a user's viewpoint. Deep forensic analysis of digital artifacts or data recovery is not the intended use of it. The homepage of Whatismyipaddress is shown in fig 5. To examine the suspect's profiles and gain knowledge about their professional backgrounds and connections with other organizations, LinkedIn is utilized. In order to support the investigation, it may involve examining user profiles, connections, posts, messages, and any other relevant data. The website can be accessed through the link https://www.linkedin.com/. The search result of background check in linkedin is showed in fig 6.

Case-2 (Insiders attack)

Te2 Technique company is a technology company that is publicly traded and known for its cutting-edge products and services. Due to upcoming product launches and strategic partnerships, the company's stock price has recently experienced significant fluctuations. Several key individuals within the organization have been implicated in a case of alleged insider trading. In the weeks leading up to a major product launch, there was a sudden surge in trading activity regarding the stock of Te2 technique company. Suspicions of insider trading are raised due to unusual fluctuations in the stock's price. An investigation was launched by investigative authorities, including the Securities and Exchange Commission (SEC).

Questions

- 1. Could the stock trading activity have been caused by unauthorized disclosures of non-public information within the company?
- 2. Did the suspicious trading activity result in any financial benefits for any of the key players?
- 3. What is the purpose of the investigation? Investigate stock market information and financial information?

Investigative tool

The census is a cybersecurity service and search engine that provides information about internet devices, domains, and certificates. This tool was utilized to monitor the resources used for wide scans and analyze various components of the internet infrastructure. Asset tracking is also a function of it. But this tool is most helpful in the case of big multinational companies. In our situation, we are employing the model of a small company, but the results obtained are not advantageous. The website can be accessed through the link https://search.censys.io/. The result obtain from census tool is showed in fig 7.Emailrep is an advanced system that gathers information about email addresses, domains, and internet personalities using crawlers, scanners, and enrichment services. The risk linked to the suspect email address was evaluated using this tool in our case. This tool allows us to comprehend email data from dark web data. Although the tool is open source, it requires payment to view the results while analyzing. By using the link





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https://emailrep.io, the website can be accessed. The website is unreliable and it takes a long time to respond. The Email rep tool's homepage can be seen in fig 8. Yahoo Finance provides a variety of services and data related to finance, investment, and the stock market, including financial headlines. The company's stock value is not present in this case. To find out if a company has experienced a sudden increase in the stock market, we can use Yahoo Finance. Insider attacks can occur for this reason, but it is not realistic if the company's stock value is not visible. The website can be accessed using the link https://finance.yahoo.com. Homepage of yahoo finance is seen in fig 9. Google Finance is a news website that offers stock quotes in real-time and delayed, interactive charts, business profiles, portfolio monitoring, economic calendars, and an industry data platform provided by Google. Regarding our case, the model company we have taken as an example does not have a stock value in Google Finance. Identifying the insider attack is not possible with this tool. Accessing the website is possible through the https://www.google.com link. The homepage of google finance is showed in fig 10.

CASE-3 (data breach)

Te2 Technique, a technology company publicly traded and renowned for its cutting-edge products and services, found itself in the midst of a significant data breach. The company's reputation was affected, and critical questions about the security of sensitive information were raised by the breach, which had wide-reaching consequences. A large amount of sensitive customer data was discovered to have been exposed by Te2 Technique. A security expert noticed unusual data traffic on the company's internal network, leading to the discovery of the breach. The presence of an unauthorized intruder who had gained access to customer data, including personal information and financial records, was confirmed through subsequent investigations.

Questions

- 1. What led the intruder to gain access to Te2 Technique's systems and compromise customer data?
- 2. What is the extent of the data breach in terms of the number of customers and records affected? Were there some high-profile clients who were impacted?
- 3. What actions were taken to contain the breach and retrieve the compromised data?

Investigative Tool

Have I been pwned? It is a website created by security researcher Troy Hunt. Assisting individuals in determining if their email addresses and associated accounts have been compromised in data breaches. In our case, we are using this tool to check whether the board of directors' email has been breached and to understand which sources it has been leaked from. Even the types of data that have been leaked in this case. The website can be accessed by using the link https://haveibeenpwned.com/. The obtained result from have I been paned tool is shown in fig 11. Zauba Corp is an Indian company that provides data on import and export transactions, company details, and other businessrelated information. To understand the company structure and details of the director related to our case, we have used this tool. It may not be the primary tool for investigating data breaches. However, it can provide us with a comprehensive understanding of the company we are collaborating with for the data breach incident. Accessing the website is possible by using the link https://www.zaubacorp.com/. The results obtained from the zauba corp website is shown in fig 12. Google Dorking, or Google hacking, is the practice of using advanced search operators to locate specific information or vulnerabilities in web applications and websites. Google Dorking can be a valuable tool when investigating a data breach to uncover publicly accessible information related to the breach or its potential sources. We can only access the company's breached data if it is available on the internet. Some versions require payment to view the results, while others are free to use. The website is accessible through the link https://www.google.com/. The example for google dork is seen in fig 13.

DISCUSSION

Data accuracy and reliability can be a significant issue, as information found in open sources may be inaccurate, outdated, or intentionally misleading, potentially leading to incorrect conclusions if not properly verified. Certain





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sources, such as subscription-based websites or proprietary databases, may be limited and lead to missing information. The verification process of online sources, especially anonymous or unverified accounts, can be challenging due to their authenticity and accuracy. In order to identify accurately, additional sources or evidence may be required because they do not have clear connections to individuals or entities responsible for business crimes.

ETHICAL CONSIDERATION

Ethical consideration is necessary for the analysis of white-collar crimes. The focus of this paper is on analyzing business crimes using the OSINT framework. We have used three different hypothesis case studies which are made up cases, and to understand how far the OSINT tools will analysis we wanted a company to run their name in different tools and identify the results. Our paper examines a service-based company called Te2 Technique that is situated in Chennai, Tamil Nadu. We have been granted permission by the company to use their company as a model.

CONCLUSION

Considering the rising number of white-collar or business-related crimes in India, obtaining a comprehensive understanding of the nature of these crimes, their methodologies, and the tools that can be used to analyze them is essential. Many digital investigative tools operate on a paid subscription basis, but in our research, we explore a range of open-source tools that are freely accessible, along with an assessment of their suitability for use in the Indian context. Data breaches, insider attacks, and money laundering are the three distinct categories of business crimes that we have focused on within this study. We used ten different Open-Source Intelligence (OSINT) tools that are readily available at no cost during our analysis of these crimes. It is important to mention that the outcomes obtained with these tools weren't always significant from an investigative perspective. Only four out of the ten tools reviewed were truly helpful in comprehending the complexity of the cases at hand. In conclusion, our findings suggest that while OSINT tools can be a valuable resource, they may not be entirely reliable for in-depth investigative purposes, and it is essential to exercise caution when interpreting the data generated by these tools, as they may contain errors.

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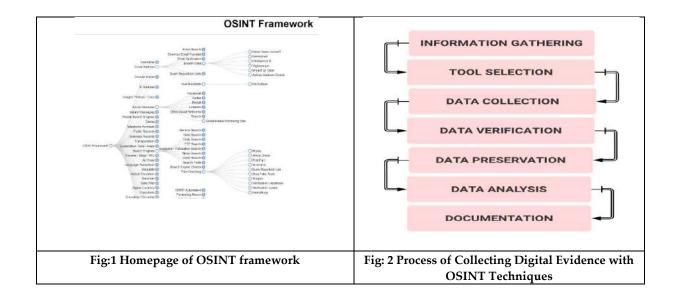
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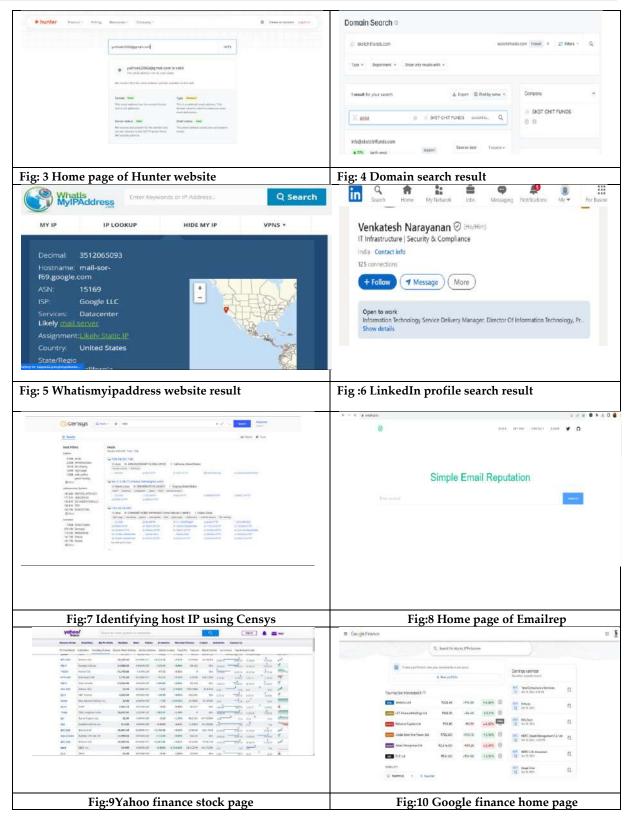
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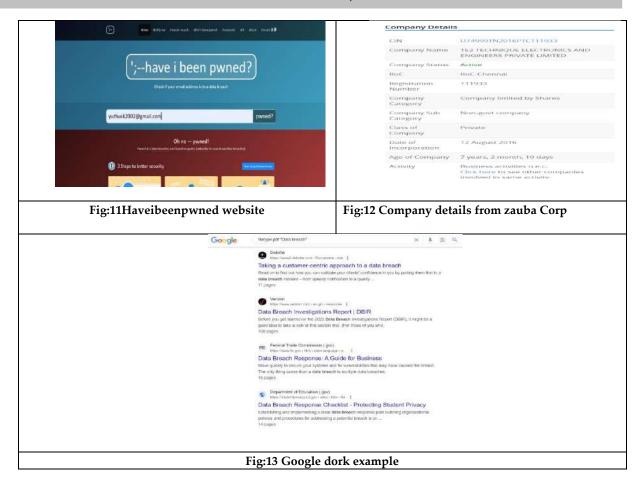
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RESEARCH ARTICLE

Cross-Cultural Management Challenges in International Business **Operations**

M.Shalini1* and M.Rohith Sundar2

¹Assistant Professor, Department of Management studies, SRM Easwari Engineering College, (Affiliated to Anna University), Chennai, Tamil Nadu, India.

²Department of Management studies, SRM Easwari Engineering College, (Affiliated to Anna University), Chennai, Tamil Nadu, India.

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*Address for Correspondence M.Shalini

Assistant Professor, Department of Management studies, SRM Easwari Engineering College, (Affiliated to Anna University), Chennai, Tamil Nadu, India. Email: Shalini.m@eec.srmrmp.edu.in,



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ABSTRACT

In today's connected world, how we communicate across cultures is becoming increasingly important, especially in international businesses affected by globalization. Managers around the world recognize the need to understand and address these challenges effectively. Globalization not only impacts our cultural identities but also shapes the way we interact with people from different backgrounds on a daily basis. To succeed in the competitive global marketplace, businesses must be equipped to anticipate and manage cultural differences. This means mastering media dynamics and international experience. By increasing awareness and experience in these areas, organizations can minimize cultural barriers and thrive in a globalized environment. Similarly, Hofstede's concept of geographical proximity may need to be reassessed in today's context of globalization. This suggests that theories of intercultural communication need to evolve, especially when applied to rapidly changing industries such as fast fashion. Our contribution goes beyond a simple case study; it challenges established theories by renowned authors, with the aim of refining and enhancing our understanding of intercultural communication in modern, globalized industries. In summary, this article highlights the importance of cross-cultural management in today's globalized business environment and provides recommendations for overcoming the challenges associated with cultural diversity. By embracing cultural differences and adopting culturally sensitive approaches, organizations can improve their competitiveness and drive sustainable growth globally.





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Keywords: Cross-cultural communication challenges, Globalization, International corporations, Managers, Global Environment, Cultural issues, International strategies, Heightened awareness, Communication dynamics.

INTRODUCTION

Cross-cultural communication challenges are increasing globally, due to the spreading influence of globalization in international business. Given the global nature of this problem, it has become an important area of exploration for managers worldwide. We acknowledge that globalization is fundamentally changing our understanding of our current culture and identity, representing a profound change in the world. In the modern context, individuals find themselves immersed in a global environment where diverse countries and cultures interact on a daily basis. Therefore, the ability to predict and manage cultural issues becomes imperative to survive the highly competitive international strategies adopted by businesses. We argue that increased awareness of communication dynamics, coupled with international experience, helps alleviate the cultural challenges inherent in doing business on a global scale. Our argument stems from the belief that a comprehensive understanding of international communication and exposure practices will significantly reduce potential cultural barriers in companies engaging in global strategy. Therefore, traditional theories of cross-cultural communication require a paradigm shift when applied to the fast fashion industry. Our contribution goes beyond a simple informative case study; it serves as a theoretical challenge to established frameworks by established authors. This suggests that the changing landscape of the fast fashion industry requires a reassessment and adjustment of existing theories of intercultural communication.

Culture

Similar to globalization, culture has evolved with many different meanings. However, for pragmatic purposes, our concept of culture is derived from a precise and clear definition. According to the Cambridge Dictionary, culture is defined as "the way of life, especially the customs and beliefs common to a particular group of people at a particular time." Looking at the historical development of the concept of culture, ethnicity emerged as a distinction between "us" and "them", in search of a common identity. It is suggested that loyalty and a non-judgmental approach should be towards one's own culture. In the context of this study, culture is considered not only from a national perspective, as described above, but also from a business or organizational perspective. Thus, in this survey, culture can be classified into three main aspects: national culture, industrial culture and corporate culture. Schein (1990) used dimensions such as internal consistency, stability, group history, learning intensity, and clarity of leader assumptions to describe organizational culture, like national culture, is diverse in nature. In addition, the growing pressures of globalization are challenging our understanding of culture and identity. Global engagement and interaction with diverse cultures involves navigating the different values, beliefs, expectations, and goals shared by members of particular groups.

Globalization

The phenomenon of globalization has penetrated even the most remote places in the world, becoming a central topic of discussion for many people. As a result, the term has taken on different meanings to different proponents. Globalization is defined as the expansion of economic activities across national borders, including international trade, investment and finance – all of which are integral components of everyday life in the world. The 21st century brings both complex opportunities and challenges. The advent of technology and advances in transportation have been key factors in solidifying globalization, regardless of consensus. In particular, the Internet presents itself as a technological masterpiece, seamlessly connecting the entire world. However, globalization also causes cultural dislocation, manifested in difficulties in adapting to new or rapidly changing cultural environments. Additionally, increased international mobility catalyzed by globalization has contributed significantly to the fast fashion phenomenon. This dynamic, influenced by increasingly global movements, encapsulates the essence of fast fashion, further highlighting the transformative impact of globalization.





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Cross-Cultural Communication

In today's connected world, globalization and cultural diversity have brought both opportunities and challenges for intercultural communication. These challenges are like obstacles that are difficult to accept and understand. Sometimes we still see a gap between "us" and "them," even in the business world, where we talk about organizational culture. As we interact across borders and engage with diverse cultures, there is no right or wrong way to do things. Everyone has their own identity, but we still need to communicate effectively to succeed in business. Good communication is essential to coordinate activities and build strong relationships, ultimately leading to better performance. In today's globalized business world, cross-cultural management is essential. Unclear messages can confuse employees and lead to negative consequences for the company. Therefore, it is important to avoid misunderstandings and ambiguity. The goal of communication is to be understood, so it's important to learn how to express yourself in a way that others can understand.

The ability to communicate effectively with diverse audiences is essential for success. Successful intercultural communication involves refraining from personal judgment, embracing diversity, showing respect, accepting ambiguity, being flexible, and being open to observing and learning. Although much research has been conducted on globalization and culture, it is important to continue to emphasize these concepts. Today, their interactions can lead to valuable business opportunities and competitive advantages. When managing cultural diversity, it is important to value differences while recognizing that individuals from one culture can learn from others and vice versa. Good intercultural communication skills are essential for the transmission of knowledge between different groups or individuals. Although cultural diversity can create communication challenges, simply recognizing the differences can help simplify the process. Good management of manners, etiquette and etiquette also contributes to successful cross-cultural business communication. This is especially true in the fast fashion space, where communication takes place in a dynamic and rapidly changing environment. Cultural differences have a significant impact on the competitiveness and performance of fast fashion companies, as the sector varies depending on the region in which they operate. Therefore, studying how cultural factors affect this industry is of great interest.

Problems

Cross-cultural management involves managing differences and similarities in values, attitudes, behaviors and work-related practices between people from different cultures. While managing diversity can bring many benefits, it also poses some challenges. Here are some of the key problems of cross-cultural management;

Communication barriers

Differences in language, cultural norms, and communication styles can lead to misunderstandings, misinterpretations, and breakdowns in communication. Direct and indirect communication styles, high and low context communication, and different nonverbal cues can all create difficulty in conveying messages accurately.

Stereotypes and biases

Cultural stereotypes and biases can lead to assumptions and biases about individuals from different cultures. These stereotypes can influence hiring, promotion, and decision-making processes, leading to discrimination and inequality in the workplace.

Conflict resolution

Cultural stereotypes and biases can lead to assumptions and biases about individuals from different cultures. These stereotypes can influence hiring, promotion, and decision-making processes, leading to discrimination and inequality in the workplace.

Leadership and management styles

Different cultures have different expectations and preferences for leadership and management styles. Authoritarian, democratic, participatory, and paternalistic leadership styles can be viewed differently depending on the cultural context. Leaders must adapt their leadership style to fit cultural preferences and expectations.





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Decision-making processes

Cultural differences in decision-making styles, such as individualism versus collectivism, hierarchical versus egalitarian decision making, short-term versus long-term orientation, can lead to challenges in decision making, consensus seeking and decision making. Understanding cultural preferences and engaging diverse perspectives can help alleviate these challenges.

Cultural adaptation and adjustment

Expatriates and multicultural groups often face challenges in adapting to new cultural environments and adjusting to cultural differences. Cultural adaptation requires flexibility, openness, empathy and intercultural skills. Lack of cultural fit can lead to stress, frustration, and decreased performance.

Ethical dilemmas

Cultural differences in ethical norms, values, and practices can create ethical dilemmas in cross-cultural business contexts. What may be considered ethical in one culture may be viewed differently in another culture. Balancing cultural relativism with moral universalism requires careful consideration and ethical decision making.

Cultural integration and cohesion

Managing diverse teams and organizations requires efforts to promote integration, engagement, and cultural inclusion. Building a shared sense of identity, promoting cross-cultural understanding and collaboration, and promoting diversity and inclusion initiatives is essential to creating multicultural teams and organizations cohesive, high performance. Overall, effectively managing cross-cultural differences requires cross-cultural awareness, sensitivity, flexibility, and competence. By proactively addressing these issues and challenges, organizations can leverage cultural diversity as a source of innovation, creativity and competitive advantage.

Measures to overcome barriers in cross-cultural communication Overcoming Language Differences and Communication Styles

To bridge the language gap and differences in communication styles, organizations should invest in language training programs and provide resources for employees to improve their language skills. Fostering a culture of open communication and encouraging questions can help clarify potential misunderstandings. Hiring an interpreter or using translation tools can also help make communication clearer.

Mitigating Stereotypes and Biases

To combat cultural stereotypes and biases, organizations must prioritize diversity and inclusion training. Implementing a blind hiring process can help eliminate bias in hiring and promotion decisions. Encouraging employees to challenge conventional wisdom and foster a culture of inclusion can contribute to a more equitable workplace.

Resolving Conflicts Effectively

Understanding different conflict management styles is important. Organizations should establish clear conflict resolution procedures, provide training on effective conflict resolution techniques, and encourage open dialogue. Creating a culture that values different perspectives and seeks common ground will foster a more collaborative and anti-conflict environment.

Adapting Leadership and Management Styles

Leaders should undergo cultural competency training to understand different leadership expectations. It is essential to adopt a flexible leadership approach that adapts to cultural preferences while preserving the organization's core values. Soliciting feedback from different team members can help leaders adapt their style to fit cultural expectations.





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Enhancing Decision-Making Processes

Fostering diversity in decision-making groups allows for a broader range of perspectives. Training on cultural differences in decision-making styles can help team members understand and appreciate different approaches. Establishing clear communication channels and encouraging transparent discussions can lead to more inclusive decision-making.

Facilitating Cultural Adaptation and Adjustment

Helping expatriates and multicultural groups adapt to new environments requires cultural orientation programs and ongoing support networks. Encouraging a culture of mutual learning, mentoring, recognition and celebration of cultural diversity within an organization can contribute to smoother transitions and increased employee satisfaction.

Navigating Ethical Dilemmas

It is important to develop a comprehensive code of ethics that takes cultural nuances into account. Ethics training should emphasize cultural relativism and moral universalism, encouraging employees to approach ethical dilemmas with cultural sensitivity. Establishing an ethics committee with diverse representation can provide guidance on complex cross-cultural ethical issues.

Promoting Cultural Integration and Cohesion

Building cultural integration includes fostering an inclusive environment where all voices are heard and valued. Creating affinity groups, promoting cross-cultural team-building activities, and celebrating cultural events can contribute to a sense of unity. Leaders must actively support diversity and inclusion initiatives to create a cohesive organizational culture.

CONCLUSION

In short, the field of international business represents a dynamic landscape where the challenges of cross-cultural communication pose both formidable obstacles and transformative opportunities. As emphasized throughout this paper, language differences, cultural nuances, patterns, and different communication styles can create complexities that affect organizational performance and efficiency. Recognizing and understanding these challenges is the first step toward building a culturally competent workforce and promoting successful international collaboration. Addressing these challenges requires a multifaceted approach, including cultural sensitivity training, leadership adaptability, conflict resolution strategies, and a commitment to diversity and get on well. As globalization continues to intertwine economies and societies, organizations that proactively address cross-cultural communication challenges not only minimize risks but also harness the power of cultural diversity such as embracing cultural differences not only improves organizational performance but also fosters innovation, creativity and adaptability on a global scale. Faced with these challenges, the call to action for international businesses is clear: invest in cultural skills, promote inclusion and foster an environment that encourages open dialogue and mutual understanding. In doing so, organizations can navigate the complexities of cross-cultural communication, foster harmonious collaboration, and achieve long-term success in the diverse and interconnected world of international business.

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RESEARCH ARTICLE

Business Analytics: Driving Sustainability through Technology of **Predictive Maintenance**

M. Shalini^{1*}, Anushya B² and Jaysre P²

¹Assistant Professor, Department of Management Studies, SRM Easwari Engineering College, (Affiliated to Anna University), Chennai, Tamil Nadu, India.

²Student, SRM Easwari Engineering College, Chennai, (Affiliated to Anna University) Tamil Nadu, India.

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*Address for Correspondence

M. Shalini

Assistant Professor, Department of Management Studies, SRM Easwari Engineering College, (Affiliated to Anna University), Chennai, Tamil Nadu, India.



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ABSTRACT

The increasing emphasis on sustainability calls for creative approaches to maximize resource efficiency. The potential of business analytics (BA) in putting predictive maintenance (PdM) ideas into practice for resource optimization is examined in this study. Maintenance schedules can be optimized and possible problems can be predicted using machine learning and sophisticated analytics algorithms that analyze large datasets on equipment performance and environmental conditions. Significant advantages come from this proactive strategy, such as decreased waste from needless replacements and repairs, longer equipment lifespans that reduce the need for manufacture and disposal, and optimized energy use by foreseeing peak demand. The paper discusses the technology, its goal, and potential solutions while presenting a framework for utilizing BA in PdM. Furthermore, real-world examples are examined to demonstrate how this approach might be used in practice. Through this integration of BA and PdM, businesses can contribute to a more sustainable future by minimizing resource usage, extending equipment life, reduce downtime and reducing their environmental impact.

Keywords: Business analytics, Predictive maintenance, machine learning, advanced analytics

INTRODUCTION

Businesses nowadays must balance reducing their environmental effect and increasing efficiency. Conventional maintenance techniques, which are frequently reactive and predicated on timetables, can result in waste production, equipment failures, and needless resource use. These actions have a detrimental impact on the entire environmental





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footprint in addition to operating expenditures. Sustainability has become a critical business imperative, demanding innovative solutions to environmental and social concerns. Businesses face the challenge of balancing economic growth with environmental responsibility and social well-being. This paper explores the transformative potential of business analytics (BA) in navigating this complex landscape. This journal investigates the use of business analytics (BA) in predictive maintenance (PdM), a proactive strategy that makes use of machine learning (ML) and data analysis to forecast equipment breakdowns and improve maintenance schedules. Businesses can reap major benefits by switching from reactive to predictive maintenance, such as:

- Decreased resource consumption
- Increased equipment lifespan
- Optimal energy consumption
- Enhanced operational efficiency

OBJECTIVES

Minimize resource consumption by using data analysis to forecast equipment breakdowns and cut down on needless maintenance and replacements. Increase the lifespan of equipment by proactively seeing any problems and planning preventative maintenance, which will cut down on the need for new equipment. Reduce energy use by anticipating peak demand and making operational adjustments based on equipment data. Boost operational effectiveness by reducing downtime with preventative maintenance to guarantee continuous operation and higher output. Improve sustainability through resource efficiency and less environmental impact, the aforementioned goals can help create a more sustainable future.

PROPOSED SOLUTION

We used actual instances from manufacturing facilities, which are crucial to the country's economy. In terms of the manufacturing company, they are fulfilling deadlines to create more, but they also need to pay attention to the environmental impact they are having. We propose that we could deploy remedies and lower downtime mistakes by doing the following actions.

DATA COLLECTION

Critical machinery has a variety of sensors attached to it that gather information on variables including temperature, vibration, pressure, and power usage. These sensors are essential for maintaining optimal operation and prompt repair by keeping an eye on the condition and performance of vital gear. Similar to a cloud platform, this data is continuously transmitted and kept in a central repository. It is simpler to examine trends, spot any problems, and make deft decisions to boost productivity and avoid unplanned downtime when data is gathered and kept in one single location. Above is a diagram of two sensors attached to the motor of a machine. This is a simple illustration to show how it works. The sensors could be temperature sensors or vibration sensors. Starting with a set of data, predictive maintenance algorithms are designed. Large data sets, including those from numerous sensors and devices operating at various times and in various environments, are frequently difficult to handle and process. Any one or more of the following categories of data may be available to you:

- Actual data from a typical system configuration
- Real information from a malfunctioning system
- Actual failure data from the system (run-to-failure data)

DATA PREPROCESSING

To guarantee the correctness and dependability of your dataset, data cleansing is essential. It entails eliminating any mistakes or discrepancies that might have been made while gathering the data. After this, feature engineering assists in extracting pertinent features from the unprocessed data. For example, it computes the temperature readings' standard deviation to capture variability. In the last steps of your analytic or modelling process, normalisation is





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used to scale the data to a common range, enabling effective comparison and study of various properties. To transform the data into a format that makes it simple to extract condition indicators, data preprocessing is frequently required. Simple methods like removing outliers and missing values are included in data preprocessing, as are more sophisticated signal processing methods like short time Fourier transforms and order domain transformations. Knowing your machine and the type of data you have can help you choose the right preprocessing techniques. For instance, choosing preprocessing approaches can be aided by understanding what frequency range is most likely to exhibit important features when filtering noisy vibration data. Transforming gearbox vibration data to the order domain, which is utilized by rotating machines when the rotational speed varies over time, may also be helpful.

MACHINE LEARNING -MODEL TRAINING MODEL SELECTION

Appropriate machine learning methods are selected based on the kind of machinery and failure prediction objectives. Popular choices include of:

- **REGRESSION MODELS** Estimate the equipment's remaining useful life.
- CLASSIFICATION MODELS Determine if the device is in good working order or in danger of breaking down.

TRAINING DATA The selected model is trained using historical data, such as sensor readings and previous failure instances. The model gains the ability to recognise trends and connections between sensor data and equipment malfunctions. The identification of condition indicators—features in your system data whose behavior varies predictably as the system deteriorates—is a crucial first step in the creation of predictive maintenance algorithms. Any feature that is helpful in differentiating between normal and malfunctioning operation or in estimating the remaining usable life might be considered a condition indicator. A helpful condition indicator separates distinct system statuses from comparable ones by grouping them together. A few instances of condition indicators are amounts obtained from

- Simple analysis, such as the mean value of the data over time
- More complex signal analysis, such as the frequency of the peak magnitude in a signal spectrum, or a statistical moment describing changes in the spectrum over time
- Model-based analysis of the data, such as the maximum eigenvalue of a state space model which has been estimated using the data
- Combination of multiple features into a single effective condition indicator (fusion)

Vibration data, for instance, can be used to track a gearbox's health. Vibrations alter in frequency and intensity when the gearbox is damaged. As a result, the peak frequency and peak magnitude serve as helpful condition indicators by revealing the type of vibrations that are occurring within the gearbox. You can derive these condition indicators by regularly analyzing vibration data in the frequency domain to keep an eye on the gearbox's health. Even when you have real or simulated data representing a range of fault conditions, you might not know how to analyze that data to identify useful condition indicators. The right condition indicators for your application depend on what type of system, system data, and system knowledge you have. Therefore, identifying condition indicators can require some trial and error, and is often iterative with the training step of the algorithm development workflow. Among the techniques commonly used for extracting condition indicators are

- Order analysis
- Modal analysis
- Spectrum analysis
- Envelope spectrum
- Fatigue analysis
- Nonlinear time-series analysis
- Model-based analysis such as residual computation, state estimation, and parameter estimation





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PREDICTION AND MAINTENANCE NEW INFORMATION & PREDICTION

The trained model is provided with real-time sensor data.

FAILURE RISK ASSESSMENT

The model makes predictions about the probability of a failure in the near future by analysing the data.

PROACTIVE MAINTENANCE

Preventing unplanned breakdowns by scheduling maintenance interventions only, when necessary, based on the projected risk. This lower:

- UNNECESSARY DOWNTIME By keeping the machinery running, production time is increased.
- **REPAIRS THAT ARE REACTIVE** Preventing issues before they arise saves money on repairs and helps to keep people safe.
- ENERGY WASTE Preventive maintenance procedures that are not necessary, such as changing out working
 parts, are kept to a minimum.

MODEL IMPROVEMENT

PERFORMANCE MONITORING

By contrasting the model's predictions with real failures, the performance of the model is continuously assessed. Retraining: To increase the model's efficacy if its accuracy starts to deteriorate over time, fresh data can be used to retrain it. Deploy and integrate the algorithm into your system once you have found a working solution for handling and processing the new system input in an appropriate manner and producing a prediction. You can use embedded devices or the cloud to implement your algorithm, depending on the details of your system. When collecting and storing a lot of data on the cloud, a cloud implementation can be helpful. The maintenance procedure is more efficient when data transfers between local computers running the prognostics and health monitoring algorithm and the cloud are eliminated. Cloud-based calculations can be made available via web apps, dashboards, email notifications, and tweets. An alternative is to have embedded devices that are closer to the equipment itself run the algorithm. The primary advantages of this approach are the reduction of data transmission volume, as information is only sent when required, and the instantaneous availability of updates and notifications regarding the condition of the equipment. Using a mix of the two is a third choice. The predictive model can operate on the cloud and send out notifications as necessary, while the preprocessing and feature extraction portions of the algorithm can be executed on embedded devices. In systems such as oil drills and aircraft engines that are run continuously and generate huge amounts of data, storing all the data on board or transmitting it is not always viable because of cellular bandwidth and cost limitations. Using an algorithm that operates on streaming data or on batches of data lets you store and send data only when needed.

COMPANIES UTILIZING SENSOR DATA AND MACHINE LEARNING FOR PREDICTIVE MAINTENANCE INDUSTRIAL TITANS

GE DIGITAL Offers predictive maintenance software and other industrial internet of things (IIoT) solutions to a range of industries, including oil and gas, aviation, and power production.

SIEMENS Provides MindSphere, an industrial open-cloud platform that helps businesses gather and evaluate sensor data for preventive maintenance.

HONEYWELL Provides hardware and software for building automation, as well as tools for equipment predictive maintenance.

TECH GIANTS

IBM: Provides the asset management Maximo Application Suite, which uses AI and machine learning to provide predictive maintenance.





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MICROSOFT Provides Azure IoT services, allowing companies to connect, monitor, and analyze data from their equipment for various purposes, including predictive maintenance.

BENEFITS OF PREDICTIVE MAINTENANCE BY BUSINESS ANALYSIS FOR SUSTAINABILITY

Businesses nowadays must balance reducing their environmental effect and increasing efficiency. Reactive maintenance techniques that are too traditional can result in waste production, unplanned equipment breakdowns, and needless resource consumption. This has an adverse effect on the entire environmental footprint in addition to operational expenditures. This problem can be solved with predictive maintenance (PdM), a proactive strategy that makes use of data analysis and business analytics (BA). Using sensor data, maintenance logs from the past, and operational data, PdM gives companies the ability to:

MINIMIZE RESOURCE CONSUMPTION

PdM reduces the requirement for needless repairs and replacements by anticipating possible equipment breakdowns and enabling targeted maintenance interventions. This results in a major decrease in the amount of energy, spare parts, and raw materials needed for reactive repairs.

INCREASED EQUIPMENT LIFESPAN

Prompt identification of possible problems enables preventative maintenance, which lowers the risk of catastrophic failures and the requirement for early equipment replacement. This lowers the environmental impact of producing and discarding new equipment in addition to saving money. Think about the energy and materials needed to produce a new industrial motor versus using PdM to make the current one last longer.

OPTIMIZE ENERGY CONSUMPTION

Businesses can predict equipment peak demand thanks to predictive analysis. Businesses can optimise energy consumption and minimise waste by making adjustments to operations and energy usage patterns based on these estimates. For instance, a plant is able to forecast the peak load that a particular machine would encounter and modify its production schedule to reduce energy consumption during such periods.

BOOST OPERATIONAL EFFICIENCY

Proactive maintenance (PdM) reduces downtime, guarantees uninterrupted operation, and boosts output. Better resource use and less waste creation across the whole production process result from this. Machine learning algorithms are able to recognise patterns in this data, create connections between possible equipment breakdowns and current operating circumstances, and eventually forecast when maintenance is required by processing and evaluating the data. This leads to tremendous resource optimization by enabling organisations to go from a reactive "fix-it-when-it-breaks" approach to a proactive "prevent-the-break" strategy. There are difficulties in putting business analytics into practise for PdM. It necessitates thorough preparation, training and technological investments, and data security assurance. Nonetheless, a more sustainable future is greatly aided by the advantages of decreased resource use, longer equipment life, optimal energy use, and increased operating efficiency. Businesses may reduce their environmental impact and guarantee long-term resource optimization for the earth and themselves by adopting this data-driven approach.

CONCLUSION

Businesses are using predictive maintenance (PdM) more and more as a potent tool for resource management and attaining sustainability objectives in response to growing environmental concerns. PdM enables a proactive approach to maintenance by utilising business analytics (BA) and the massive amount of data gathered from sensors





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implanted into equipment. This method anticipates possible equipment failures and moves away from reactive repairs based on breakdowns. There are numerous advantages to this proactive approach for businesses and the environment. PdM reduces resource use, to start. The demand for raw materials, spare parts, and the energy required for reactive repairs is decreased when equipment data is analysed to forecast and prevent needless repairs and replacements. Furthermore, PdM increases equipment longevity by facilitating timely preventative measures. As a result, there is a decrease in the frequency of equipment changes, which lessens the environmental effect of producing and discarding new equipment. Optimized energy use is yet another important advantage. Businesses can reduce energy use and carbon emissions by modifying operations and energy usage patterns in response to anticipated equipment peak demand. In general, firms are empowered to function more sustainably when BA and PdM are integrated. This strategy reduces environmental effect, promotes resource efficiency, and advances a more sustainable future. PdM has the potential to become a pillar of sustainable business practises for companies in all sectors as this technology develops and becomes more widely available.

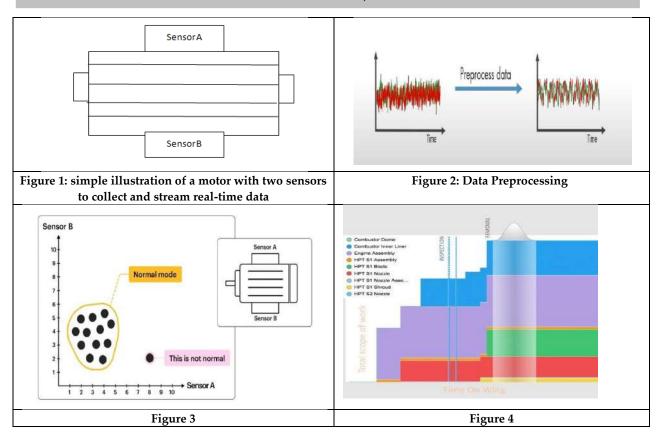
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RESEARCH ARTICLE

Adapting to the Dynamic Digital Marketing Landscape: Trends and **Strategies for Success**

Saikumari. V^{1*}, Prathosh K.D², Priyanka R² and Ramya Amalraj S²

¹Professor and Head, MBA Department, SRM Easwari Engineering College, Chennai (Affiliated to Anna University) Tamil Nadu, India.

²Student, Department of Master of Business Administration, SRM Easwari Engineering College, Chennai (Affiliated to Anna University) Tamil Nadu, India.

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*Address for Correspondence Saikumari. V

Professor and Head, MBA Department, SRM Easwari Engineering College, Chennai (Affiliated to Anna University) Tamil Nadu, India.



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ABSTRACT

The digital marketing landscape is a dynamic tapestry, constantly evolving due to technological advancements, shifting consumer behaviours, and the ever-changing nature of global communication. This comprehensive review delves into recent advancements and strategic trends shaping the future of the field, focusing on key areas: artificial intelligence (AI), the dominance of short-form video content, the primacy of customer experience (CX), the evolution of influencer marketing, and the potential of the metaverse. It emphasizes the crucial role of remaining agile and embracing a continuous learning mindset to excel in this ever-shifting digital environment.

Keywords: Digital marketing trends, Innovation and Evolution, Strategic focus, Expanded landscape.

INTRODUCTION

The digital revolution has fundamentally transformed how businesses interact with their target audiences. Digital marketing has become an indispensable tool, enabling brands to reach a global audience, build relationships, and drive conversions. However, the digital landscape is not static. It is constantly evolving, propelled by technological innovation, evolving consumer behaviour, and the emergence of new trends and platforms. This paper provides a comprehensive exploration of this dynamic landscape. It examines recent advancements and strategic trends that are shaping the future of digital marketing and underscores the importance of continuous learning and adaptation for success in this ever-changing environment. By understanding the latest developments and embracing a strategic





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approach, businesses can leverage the power of digital marketing to achieve their goals and thrive in this dynamic environment.

KEY AREAS OF INNOVATION AND EVOLUTION ARTIFICIAL INTELLIGENCE REVOLUTIONIZES MARKETING

AI is rapidly transforming various aspects of digital marketing, offering increased personalization, automation, and predictive capabilities.

- **PERSONALIZATION:** AI algorithms analyse customer data to personalize content, recommendations, and advertising, enhancing the relevance and effectiveness of marketing campaigns [1].
- AUTOMATION: AI-powered tools automate repetitive tasks like ad bidding, social media scheduling, and content curation, freeing up marketers for strategic endeavours [2].
- **PREDICTIVE ANALYTICS:** AI can predict customer behaviour and preferences, allowing marketers to proactively deliver targeted campaigns and optimize marketing efforts for maximum impact [3].

SHORT-FORM VIDEO CONTENT TAKES CENTER STAGE

Platforms like TikTok and Instagram Reels have propelled short-form video content to the forefront of digital marketing strategies, offering unique advantages:

- ENGAGING AND ATTENTION-GRABBING: Short-form videos capture attention with their concise and visually appealing nature, making them ideal for conveying information within a limited timeframe [4].
- ADAPTABILITY ACROSS PLATFORMS: This content format is easily adaptable for various platforms, maximizing reach and audience engagement [5].
- STORYTELLING POTENTIAL: Short-form videos can be powerful storytelling tools, building brand awareness and fostering emotional connections with audiences [6].

CUSTOMER EXPERIENCE REIGNS SUPREME

Prioritizing customer experience (CX) is essential for digital marketing success. This entails focusing on:

- **PERSONALIZED INTERACTIONS:** Tailored communication and interactions build loyalty and a positive brand perception [7].
- OMNICHANNEL ENGAGEMENT: Providing a seamless and consistent brand experience across multiple touchpoints is vital for a cohesive customer journey [8].
- **PROACTIVE CUSTOMER SERVICE:** Anticipating customer needs and offering proactive support builds trust and drives brand advocacy [9].

INFLUENCER MARKETING CONTINUES TO EVOLVE

Influencer marketing remains a powerful tool but is evolving with notable trends:

- RISE OF MICRO-INFLUENCERS: Consumers are increasingly drawn to micro-influencers for their genuine connections and recommendations within niche communities [10].
- FOCUS ON AUTHENTICITY AND TRANSPARENCY: Authenticity and transparency from influencers are paramount, requiring brand partnerships to align organically with the influencer's values [11].
- LIVE SHOPPING AND INTERACTIVE CONTENT: Live streaming platforms are being leveraged for interactive shopping experiences, further enhancing influencer-audience engagement [12].

EXPANDING THE DIGITAL MARKETING LANDSCAPE THE RISE OF THE METAVERSE

The metaverse, a burgeoning virtual world experience, presents potential opportunities for marketing, such as:

• IMMERSIVE ADVERTISING EXPERIENCES: Engaging consumers in multi-dimensional marketing campaigns, blurring lines between reality and digital content.





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- VIRTUAL PRODUCT LAUNCHES: Unveiling new products in simulated environments, allowing for interactive exploration and global access.
- **INTERACTIVE COMMUNITIES:** Building online spaces for brands and consumers to connect, fostering engagement and fostering a sense of belonging.

THE CONTINUED IMPORTANCE OF DATA PRIVACY

With heightened public awareness and stricter regulations surrounding data privacy, marketers need to prioritize responsible data collection and usage. This includes:

- **IMPLEMENTING TRANSPARENT DATA PRACTICES:** Informing users about what data is collected, how it's used, and with whom it's shared, fostering trust and understanding.
- **OBTAINING EXPLICIT USER CONSENT:** Clearly requesting and receiving permission from users before collecting or using their data, ensuring respect for their privacy.
- INVESTING IN DATA SECURITY: Implementing robust measures like encryption and access controls to protect user data from unauthorized access, breaches, or misuse.

THE CONTINUED IMPORTANCE OF DATA PRIVACY

With heightened public awareness and stricter regulations surrounding data privacy, marketers need to prioritize responsible data collection and usage. This includes:

- **Implementing transparent data practices:** Clearly explain what data you collect, how it is used, and user rights regarding their data.
- **Obtaining explicit user consent:** Be upfront about data collection and obtain explicit consent before gathering any personal information.
- Investing in data security: Implement robust security measures to protect user data from unauthorized access
 or breaches.

THE INTEGRATION OF AUGMENTED REALITY (AR) AND VIRTUAL REALITY (VR)

AR and VR technologies are finding increasing applications in marketing campaigns, offering:

- **INTERACTIVE PRODUCT EXPERIENCES:** Allow customers to virtually try on clothes, see furniture in their homes, or experience travel destinations.
- VIRTUAL STORE TOURS: Create immersive and interactive virtual tours of physical stores, allowing customers to explore products and layouts from anywhere.
- LOCATION-BASED MARKETING: Deliver targeted marketing messages and promotions based on a user's location through AR technology.

THE POWER OF USER-GENERATED CONTENT (UGC):

User-generated content, such as reviews, social media posts, and customer testimonials, holds significant power in influencing consumer behaviour. Marketers can leverage UGC by:

- ENCOURAGING CUSTOMER PARTICIPATION: Run contests, giveaways, and encourage customer reviews to generate content.
- SHOWCASING POSITIVE EXPERIENCES: Feature positive customer testimonials and reviews on your website and marketing materials.
- FOSTERING BRAND ADVOCACY: Partner with engaged customers to create user-generated content and build brand loyalty.

THE EVOLVING LANDSCAPE OF SOCIAL COMMERCE

The ability to seamlessly purchase products directly within social media platforms is changing the way consumers shop. Marketers need to adapt their strategies by:

• INTEGRATING SOCIAL COMMERCE FEATURES: Explore and leverage social commerce features offered by platforms like Instagram Shopping and Facebook Shops [15].





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- OFFERING SMOOTH CUSTOMER JOURNEYS: Ensure a seamless purchasing experience within the social media platform, minimizing friction and maximizing conversion rates [16].
- LEVERAGING THE POWER OF INFLUENCERS: Collaborate with influencers to showcase products and drive sales through social commerce channels [17].

THE GROWING FOCUS ON SUSTAINABILITY:

Consumers are increasingly conscious of the environmental impact of their choices, making sustainability a significant consideration in marketing strategies. This includes:

- HIGHLIGHTING SUSTAINABLE PRACTICES: Showcase your commitment to sustainability throughout your supply chain and operations.
- OFFERING ECO-FRIENDLY PRODUCTS: Develop and market eco-friendly products that align with consumer values.
- **COMMUNICATING TRANSPARENTLY:** Be transparent about your sustainability efforts and avoid greenwashing practices.

THE NEED FOR CONTINUOUS LEARNING AND ADAPTATION

The digital marketing landscape is constantly evolving, demanding continuous learning and adaptation from professionals. This includes:

- STAYING INFORMED ABOUT EMERGING TRENDS: Attend industry events, subscribe to industry publications, and actively seek out new knowledge.
- UPSKILLING THROUGH COURSES AND WORKSHOPS: Invest in your professional development by taking courses and workshops relevant to the latest trends and technologies.
- **EXPERIMENTING AND ANALYSING RESULTS:** Don't be afraid to experiment with new strategies and analyse the results to understand what works best for your target audience.

GLOBAL IMPACT OF DIGITAL MARKETING

The internet has exploded in usage over the past few decades, as shown in the graph below. This surge in internet users has created a massive global audience for digital marketing efforts. As of October 2023, there are over 4.7 billion social media users globally, highlighting the power of this platform for brand awareness and engagement [13].

DIGITAL MARKETING'S INFLUENCE CAN BE SEEN IN VARIOUS ASPECTS OF OUR LIVES:

- **E-COMMERCE:** The ability to purchase products online has transformed the retail landscape, with global ecommerce sales projected to reach \$6.7 trillion in 2023 [14].
- SOCIAL MEDIA INTEGRATION: Social media platforms have become ingrained in our daily lives, influencing purchasing decisions, entertainment consumption, and news sources.
- MOBILE MARKETING: The increasing use of smartphones has led to the rise of mobile marketing, with businesses reaching customers through SMS, app notifications, and social media on their mobile devices.

THE FUTURE OF DIGITAL MARKETING

As technology continues to evolve and consumer behaviour shifts, digital marketing will undoubtedly continue to transform. Staying ahead of the curve by embracing new technologies, adapting strategies, and prioritizing ethical practices will be crucial for success in the ever-changing digital landscape.

CONCLUSION

The digital marketing landscape is a dynamic and ever-evolving ecosystem. Understanding its evolution and emerging trends, while upholding ethical practices and focusing on continuous learning, is essential for navigating this evolving landscape and achieving long-term strategic success. By embracing innovation, adapting strategies, and





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prioritizing ethical considerations, marketers can leverage the power of digital marketing to connect with global audiences, build meaningful relationships, and drive sustainable growth in the years to come.

THIS COMPREHENSIVE REVIEW HAS EXPLORED KEY AREAS OF INNOVATION AND EVOLUTION IN DIGITAL MARKETING, HIGHLIGHTING THE IMPORTANCE OF:

- Artificial intelligence (AI) for personalization, automation, and predictive analytics.
- Short-form video content for engaging storytelling and audience capture.
- Customer experience (CX) for building loyalty and fostering positive brand perception.
- Influencer marketing with a focus on authenticity, transparency, and micro-influencers.
- Emerging trends like the metaverse, voice search, data privacy, AR/VR, UGC, social commerce, and sustainability.
- Continuous learning and adaptation to stay ahead of the curve in this dynamic field.

By implementing these insights and embracing a strategic approach, businesses can leverage the power of digital marketing to thrive in the ever-changing digital world.

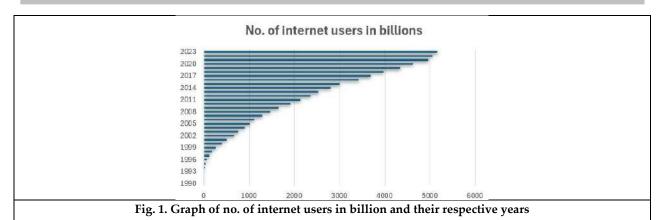
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RESEARCH ARTICLE

A Study on the Influence of Digital Finance on Individual Behaviour towards Financial Investments

Dhanya Nair*

Research Scholar, Hindusthan College of Arts and Science, (Affiliated to Bharathiyar University) Coimbatore, Tamil Nadu, India.

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*Address for Correspondence Dhanya Nair

Research Scholar, Hindusthan College of Arts and Science, (Affiliated to Bharathiyar University) Coimbatore, Tamil Nadu, India.



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ABSTRACT

Digital finance is the term used to describe the impact of new technologies on the financial services industry. It includes variety of products, applications, processes and business models that have transformed the traditional way of providing banking and financial services. The term financial technology was first coined in 1990s to refer to financial services available online but were later referred to as digital finance or e-finance in the 2000s. The Covid-19 provided additional support for the banking sector's adoption of digitization. Digital financial services, namely digital wallets, made it possible to deliver money to people in need. Consumers today need a simple, affordable solution for accessing financial services. Consumers are shifting toward digital finance as a result of the younger generation's increased propensity to use technology, including the internet and digital platforms. The digital revolution in the Indian economy will lead to a spin off effect on every industry. The adoption of new and emerging technologies has taken a center stage providing consumers with a variety of digital offering to attract them towards its usage. Amidst this revolution, the finance industry moved forward with the change and introduced digital finance. Digital finance is the root towards attaining economic development. Digital finance is the lifeline for economic enhancement enabling businesses, and individuals to invest, save, and pay through digital platforms. The present study is undertaken to highlight the usage of digital technologies in the investment pattern of individuals of Federal Bank in Kottayam dt. The study will benefit the bank to find out the awareness, usage and satisfaction level of customers in using digital solution for investment and can take decisions in modification of any of the applications currently used.

Keywords: Digital Finance, Digitization, Awareness, Usage, Satisfaction





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INTRODUCTION

Digital Financial Services (DFS) include a broad range of financial services accessed and delivered through digital channels, including payments, credit, savings, remittances and insurance. Digital channels refers to the internet, mobile phones, ATMs, POS terminals etc. Everything is going digital and so are financial services. Gone are the days when you had to carry money when going out. A mobile phone is sufficient to take care of all your daily financial needs. All banking services are within your palm. There is no need to go to a bank to get your transactions done. You can deposit, transfer, withdraw or invest money using digital devices. Digital financial services have expanded greatly in India, and people have accepted them withopen arms. Their convenience and safety prompt more people to move to these methods.

DIGITAL BANKING

A digital bank represents a virtual process that includes online banking, mobile banking, and beyond. As an end-to-end platform, digital banking must encompass the front end that consumers see, the back end that bankers see through their servers and admin control panels, and the middleware that connects these nodes. Ultimately, a digital bank should facilitate all functional levels of banking on all service delivery platforms. In other words, it should have all the same functions as a head office, branch office, online service, bank cards, ATMs, and point- of-sale (POS) machines. The reason digital banking is more than just a mobile or online platform is that it includes middleware solutions. Middleware is software that bridges operating systems or databases with other applications. Financial industry departments such as risk management, product development, and marketing must also be included in the middle and back ends to truly be considered a complete digital bank. Financial institutions must be at the forefront of the latest technology to ensure security and compliance with government regulations.

DIFFERENT TYPES OF DIGITAL BANKING SERVICES

UPI (UNIFIED PAYMENT INTERFACE)

Among the popular types of digital payments, the most popular is UPI. This allows money transfer from your bank account using a single window directly to the vendor from your mobile. The payee's virtual address with consent for mobile payment needs to be entered for this mode of digital payment. Several bank accounts can be linked with one app.

INTERNET BANKING

Internet banking or net banking is among the oldest types of digital banking. You can manageyour bank accounts virtually if you have an internet connection. Customers can visit the bank portal and enter login details (password and username) and register. The bank takes layers of security measures into account, ensuring caution during payment and transactions.

MOBILE BANKING

Another popular type of digital banking is <u>mobile banking</u>. Account holders download the bankapplication on their mobile. All bank services like balance enquiries, payments etc. can be accessed from smartphones.

BANKING CARDS

Banking cards are essential for all types of digital payments. Cards are based on their issuance, usage, and modes of digital payment. Four types of banking cards are Debit, Credit, Prepaid and Electronic cards.

- 1. Debit cards issued by the bank are linked to the bank account, and cash can be withdrawn from the ATM, this banking card allows virtual payment for products and services bought through eCommerce or offline stores.
- 2. Credit cards are issued by banks/ non-banks. These prepaid cards are used for buying products and services on credit.





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- 3. Bank Prepaid Cards are not linked to bank accounts and aid overdraft facilities. These cards need preloaded value addition and only the amount added can be spent.
- 4. Virtual Debit Card are electronic cards that work like debit cards are used for online ecommerce transactions.

MOBILE WALLETS

Mobile wallets are the digital version of a wallet that enables the user to make different types of digital payments online. Digital wallet stores money added by the user linked to their bank account. Safe amongst all digital payment methods, this can be used for all transactions throughan app installed on the smartphone and an internet connection.

BHARAT INTERFACE FOR MONEY

Bharat Interface for Money (BHIM) app enables simple payment transactions amongst other modes of digital payment. With the Unified Payments Interface (UPI) you can make direct and instant bank to bank payments and collect money using your mobile number and address. You can make direct bank payments to anyone on UPI using their UPI ID or scanning their QR withthe BHIM app. You can also request money through the app from a UPI ID. **POINT of SALE (PoS).** This is an internet linked electronic swipe machine through which a Merchant Establishment (ME) performs a retail transaction by swiping customer debit and credit card. The merchant calculates the amount owed by the customer. After receiving the amount, the merchant may issue a printed receipt or send an electronic receipt to the customer. Digital Financial Services Used in Federal Bank for Investment

Purpose

- 1. FedMobile: This is a mobile banking application used for all sorts of deposits, mutual funds, sovereign gold bond investment, TD, insurance linked products etc.
- 2. FedNet: This is a net banking facility used for all sorts of deposits and investments, mutual funds, ASBA(IPO), insurance linked products etc
- 3. Fed-E-Trade: Used for investment in shares online.

BENEFITS OF DIGITAL BANKING

CONVENIENCE

The ability to bank wherever and whenever you want is one of the main benefits of mobile and online banking solutions. Many mobile banking apps, for instance, let you deposit checks remotely. At the same time, you can check your balance, transfer funds and set up a notification to alert you if you overdraftyour account—all without the need to visit a branch. It's a real time-saver. Digital banking also offers additional conveniences, such as the ability to go cashless. Paying with cash isn't as convenient as an electronic transaction. Electronic transactions are more secure (you aren't carrying cash), they're more sanitary (you aren't touching cash) and you cantrack your transaction electronically. A cashless society with digital transactions is much more efficient, and it allows for much better management of your financial resources.

FEATURES

Many banks' mobile and online experiences offer just as many features as banking in person—if not more. Banks might offer personalized financial advice, savings tools, big-purchase calculators or even virtual assistants, all within the convenience of an app. Banking apps typically let you complete everyday banking tasks, like viewing statements and account balances, transferring funds and paying bills. Mobile check deposit, which lets users cash checks from their phone, is also common. Features like peer-to-peer payments might not be top-of-mind, but the ability to send money in minutes through your mobile banking app canbe handy, and many banks now offer this feature. Locating nearby ATMs, cardless ATM withdrawals and budgeting and tracking tools are other perks your mobile banking app may offer. Some banks even offer the ability to chat with a live representative through their apps, which can bridge the gap between in-person and digital banking. Consumers should seek out banks that prioritize offering a human touch even in their digital channels, striking the right balance between the human element and digital automation.





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SECURITY

Security is a priority for financial institutions, and that extends to mobile and online banking. Threats exist everywhere, including inside bank branches. Fortunately, many banks make it easy to take extra security precautions. For example, your bank may let you add multifactor authentication to your mobile app and online bank account. Many mobile banking apps also let you use biometric authentication to log in. Federal Bank's app, for instance, provides biometric login options that require your fingerprint or facial recognition. Your bank may also scan for certain risks automatically. Banks asks for additional verification if it detects a login from an unknown device. Overall, you may be more secure thanyou think when using digital banking. It's been reported that digital payments and e-wallets actually offer more security in some cases than a physical card, giving some users even more reason to use digital banking tools.

CONTROL

Having control over your finances with the ability to self-serve is another significant benefit of digital banking, as is real-time access to managing and moving money. Unlike banking in person, mobile banking apps and websites generally have no restrictions on when you can perform banking tasks, like depositing a cheque or moving money from one account to another. And it's getting easier to navigate daily transactions. The world of technology is offering the opportunity to be able to receive money and to spend money in ways that are much easier than they were in past times. Banks are continuing to advance the features offered on their digital banking platforms. Automated savings tools and push notifications for events like low balances or overdrafts are commonplace. In many cases, you can even activate a new debit or credit card from your app.

BENEFITS BEYOND BANKING

On a larger scale, we can use modern connectivity tools to create financial, social and economic change. This online banking product allows for a broader base of communication that can be used for things like teaching financial literacy. Digital banking is also becoming a way to find communities and options tailored to your needs as a banking customer.

CONTENT

OBJECTIVES

- 1. To know the awareness level of the individuals regarding the applications used for investment purpose in Federal Bank.
- 2. To assess the satisfaction level of the individuals by using the applications for the investment purpose in Federal Bank.
- 3. To know in which of the investment avenues the individuals invest using the applications in Federal Bank.
- 4. To know the reasons of such individual behaviour based on the applications used for investment purpose in Federal Bank.

RESEARCH METHODOLOGY

The present research design adopted in the study was descriptive nature. The study is based on primary data and secondary data. The data has been collected from individuals through the questionnaire and interview. The sampling technique used in this study is random sampling. As per Cochran, the formula for unknown population to collect the samples is:

Sample size(N) - $Z^2 PQ$

2

Where N is the sample size, e is the margin of error (5%), p is the confidence level (95%), and z is the z-score (1.96). According to this formulae, the sample size was taken as 385 participants minimum. Here data has been collected from 410 respondents through questionnaire.





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LITERATURE REVIEW

An evaluation of digital banking in Kerala with special reference to its awareness, adoption, satisfaction and loyalty by Devikrishna V(2022). The study focused on the awareness, adoption, satisfaction and loyalty of the customers towards digital banking products and services. It is inferred from the study that there is high level awareness of internet banking products and services and the customers are using those services but the customers are sometimes using all other digital banking services like ATM, phone banking and mobile banking. Influence of behavioural factors on the adoption of digital finance, an empirical study by Jain Niyati (2022). This study focused on understanding human behaviour and perception towards digital finance. For this behavioural factors have been divided into 3 aspects-cognitive, theory of planned behaviour and behavioural biases. It was inferred that most of the individuals was aware of the digital financesolution. The mostly used digital financial services were debit/credit cards followed by e-wallets, mobile/online trading. Influence of customers digital persona on their online buying behavioural aspects by Rahul Shandilya(2022). The focus of the study is to understand the personality traits which has a favourable impact on the individuals digital quotient. The personality traits were denoted by UCCCEEE (updated, confident, curious, efficient, experimentative, epicurean). The researcher fund out that the personality traits have a positive correlation with the human behaviour.

RESEARCH GAP

The above reviews shows that many researches has been conducted on digital finance. Theresearches shows the impact of digital finance in the economy, consumers behavioral pattern towards digital solution, impact on financial inclusion etc. There are no studies which has made an attempt to find out the behaviors of individuals by using digital technologies in banks only for the purpose of investment purpose. As investment is a great source of passive income and it creates a greater pay off in the future than what was originally put in, it is important and using in studying the individual behavior/attitude of individuals in using digital technologies only forinvestment purpose.

DATA ANALYSIS AND INTERPRETATION

CHI-SQUARE TEST

H0: There is no significant relation between literacy level and usage of digital applications (exclusively for investment) in Federal Bank.

H1: There is significant relation between literacy level and usage of digital applications(exclusively for investment) in Federal Bank.

CV = 29.25

TB = 15.507

DF = (r-1)(c-1) ie (5-1)(3-1) = 4*2 = 8

Analysis

As per the result calculated, it shows that there is significant relation between literacy level and the usage of digital applications. Since the calculated value is 29.25 and table value with eight degree of freedom is 15.507(0.05% of significance level), we reject H0.

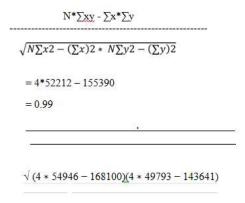
CORRELATION

Correlation between income and usage of application.





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ANALYSIS

As per the correlation data received, there is high positive correlation between the income level of customers and usage of applications. This shows that as the income level increases, people uses more applications of Federal Bank for the purpose of investment.

PERCENTAGE ANALYSIS

Analysis

As per the above data, it shows that there is a strong need for digital finance technologies in bank for investment. As per the customers opinion the applications are user friendly, leads to safety of customers, helps in saving time and cost and finally it replaces traditional banking and all the issues related to traditional banking.

FINDINGS

- 1. The chi-square test calculated gives the value of 29.25 which is more than the table value i.e 15.507 which shows that there is significant relation between literacy leveland usage of digital applications.
- 2. There is high positive correlation between the income and usage of application. As the income of the customers increases, their usage of digital applications also increases.
- 3. The data collected from the customers shows that there is a strong need for digital finance technology in the bank for investment.
- The customers says that digital finance technology leads to time and cost saving. It tends to safety of customers.
- 5. The customers also opines that the digital technologies are user friendly. It replacestraditional banking.

SUGGESTIONS

- Even though the majority of customers are satisfied with the applications there are somecustomers who are not satisfied or less satisfied. The banking officials must identify the dissatisfied customers and find a solution for that
- 2. The banking officials must continuously conduct a review from the customers to knowthe feedback.
- 3. While introducing any new applications in the bank, proper training should be provided to the employees by the top management.
- 4. Simple and easily handled applications should be introduced so that even though an illiterate or less literate individuals could be used.





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CONCLUSION

Finding out the influence of Digital Finance on individuals behaviour towards financial investment in Federal Bank with reference to Kottayam dt was the primary aim of the research. The study shows that 50.49% of customers says that there is strong need for Digital Finance technology in bank for investment. The studies also revealed that 74% of customers says that the applications are user friendly. Around 50% of customers says that it replaces traditional banking. Even though most of the customers are satisfied with the application, there are dissatisfied customers too. The banking officials must identify the dissatisfied and find a solution for that. The officials must continuously conduct a review from the customers to knowthe feedback.

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Table 1: Chi-square test between literacy level and usage of applications

	Literacy Level						
		10th	12th	UG	PG	Others	Total
	FedMobile	7	8	72	153	4	244
	FedNet	5	7	15	113	1	141
Usage	Fed-e- Trade	1	-	12	12	1	25
		12	15	99	278	6	410

(Source: Online survey, January 2024)

Table 2: Calculated Values of Chi-square test

Observed Values (O)	Expected Values (E)	$\sum (\mathbf{O} - \mathbf{E})^2$	Σ(O-E) ² /E
7	7.14	0.0196	0.0027
8	8.93	0.8649	0.097
72	58.92	171.08	2.90
153	165.44	154.75	0.94
4	3.57	0.1849	0.05
5	4.13	0.7569	0.18
7	5.16	3.3856	0.66
15	34.05	362.90	10.66
113	95.60	302.76	3.17
1	2.06	1.1236	0.55
-	0.73	0.5329	0.73
-	0.91	0.8281	0.91
12	6.04	35.5216	5.88
12	16.95	24.5025	1.45
1	0.37	0.3969	1.07
		Total	29.25

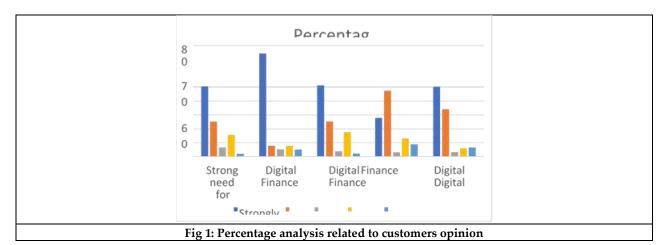


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Table 3: Correlation between income and usage of application

Range	x (Income)	y (Usage of applications)	x ²	y2	ху
Below 2 lakh	25	14	62 5	196	350
2 – 4 lakh	184	180	33856	32400	33120
4 – 6 lakh	89	86	7921	7396	7654
Above 6 lakh	112	99	12544	9801	11088
	410	379	54946	49793	52212

(Source: Online survey, January 2024)







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RESEARCH ARTICLE

The Impact of Remote Work on Employee Engagement: Strategies for **Building Connection in a Virtual Environment**

Tiya Thomas^{1*} and S.Annie Priyadharshini²

¹Research Scholar, Karunya Institute of Technology and Sciences, (Affiliated to Bharathiyar University) Coimbatore, Tamil Nadu, India.

²Assistant professor, Karunya Institute of Technology and Sciences, Coimbatore (Affiliated to Bharathiyar University) Tamil Nadu, India.

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*Address for Correspondence

Tiya Thomas

Research Scholar, Karunya Institute of Technology and Sciences, (Affiliated to Bharathiyar University) Coimbatore, Tamil Nadu, India.



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ABSTRACT

With the shift to remote work, organizations are facing new challenges in keeping employees engaged. Employee engagement is a vital notion in the attempt to comprehend and define, both qualitatively and quantitatively, the nature of the connection between an organization and its employees. This paper could look at how remote work affects employee engagement, taking into account factors like communication, collaboration, job satisfaction, and more. It could also look at strategies and best practices that organizations can implement to help their remote employees feel connected, purposeful, and engaged. The purpose of this paper is that it could help create a better virtual work environment, and also aim to discuss the psychological and emotional aspects of remote work, considering factors like work-life balance, motivation, and feelings of isolation. The study is based on the IT employees at Info park and utilizes primary data and secondary data and chi-square test for the study and has followed descriptive analysis. The findings would be to provide insights into effective strategies that organizations can implement to ensure high levels of employee engagement in a virtual work environment. This paper examines an in-depth exploration of the challenges of remote work to maintain and boost employee engagement in a virtual setting.

Keywords: Communication, Job satisfaction, strategies, psychological and emotional aspects, remote work, virtual work environment





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INTRODUCTION

Employee engagement is traditionally defined as a sense of loyalty, commitment, and employees' mental and emotional bonds with their workplaces. An individual is more likely to show increased productivity, job satisfaction, and willingness to go above and beyond in their work when they are engaged. The introduction of computers during the 1970 oil crisis marked the beginning of the "telecommuting" trend for white-collar occupations. However, as computers advanced in the 1980s, executives and upper managers began adopting work from home policies. Fast communication via fax and phone has led to an increase in work from home jobs. Additionally, the US government gave incentives to businesses that allowed their physically challenged employees to work from home. The objective is to enhance an employee's emotional attachment to the company, their role within the organization, their peer environment, and the company culture. There's been an increase in remote employment for years, and this trend is only going to continue. A growing number of businesses and workers are seeing the benefits of working remotely as they adapt to this new distant norm. It's typical for remote workers to be responsible for their own software and hardware: The majority of remote workers must pay for their devices, internet, and additional software in order to finish their work, even if some businesses provide laptops, internet, and software tools needed for remote work.

RESEARCH METHODOLOGY

There are two sources of collecting data: - (a) Primary sources and (b) Secondary sources. To collect information related to employee engagement and remote work and using various strategies in virtual environment. The Primary data was collected by distributing questionnaires via Google forms and collected the responses. The questionnaire was structured. The questionnaire contained general questions which consisted of the demographic variables such as age, gender, occupation, job role and so on. Specific questions were included which were related to the topic under study. The data was collected from 117 respondents and percentage analysis was used to conduct the data analysis. The data collected was tabulated and charts like column chart, bar charts, were used to interpret the data.

PRIMARY DATA

The primary data is primarily gathered through the distribution of questionnaires. Primary data are genuine facts that were acquired especially for the intended purpose are known as primary data. It is beneficial for future research as well as on-going study. The primary data is collected from IT professionals, Info park ,Kochi.

DATA ANALYSIS AND INTERPRETATION

Interpretation: From the above table it is clear that ;out of 117 respondents and 24.8% of the respondents belong to the age group of 18-24years, 39.3% of the respondents belong to 25-34 years, 23.1% belong to 35-44,10.3% belong to 45-54years and only 2.6% are above the age group of 55 or above.

Interpretation: From the above table we concluded that only 117 responses received on gender category and 41% of the respondents are male and 44.2% of the respondents are female, other category is not responded.

Interpretation: The above table explains the remote work period of the 117 respondents and

28.2% worked for less than one year,46.2% worked for a period between 1-2 years and 15.4% worked for 3-5 years, 10.3% worked above 5 years.

Interpretation: The above table explains Job role of remote employees. From 117 respondents,29.1% consist of Frontline employees,42.7% does middle management,12.8% belong to senior management and 15.4% with other job roles.

SECONDARY DATA

Secondary data is collected using Google scholar, books, websites, news articles. From the above statistical diagram, there will be an immense increase in the global digital jobs. The numbers of jobs (in millions) increasing by years and by 2030 remote digital jobs will rise from 25% to 92 million.





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STATISTICAL TEST

Influence of gender on remote job satisfaction

H0: There is no influence of gender on remote job satisfaction.

H1: There is significant influence of gender on remote job satisfaction.

From Table 5 it is interpreted that gender have no influence on the remote job satisfaction Here H0 is accepted as P value is more than 0.05 and H1 is rejected. There is no significant relationship between gender and remote job satisfaction.

Impact of Age on Job role of employees

H0: Age has no impact on employee job role.

H1: Age has a significant impact on employee job role.

From Table 6 it is interpreted that Age has a significant impact on employee job role. Here H0 is rejected as P value is less than 0.05 and H1 is accepted. So there is highly significant impact of Age on employee job role. There is a relationship between Age and employee job role.

REVIEW OF LITERATURE

Adisa et al. (2021)states that the during the pandemic, the abrupt shift from traditional in-person to virtual work environments resulted in increased workloads, a rise in online presenters, job insecurity, and a lack of adaptability to remote work arrangements. These stressors have the potential to drain important personal and social resources, which will have a detrimental effect on employee engagement levels. Anand et al. (2021) Points at two aspects of telecommuting: a. In normal times b. In times of crisis and in both scenarios, how do you keep remote workers engaged and motivated to ensure minimal, if not zero, loss of productivity. Examine several aspects of the remote worker's character and offers suggestions and methods to improve remote workers' involvement and explores different facets of the remote worker's persona and provides recommendations and practices to enhance employee engagement of the employees working remotely. De-La-Calle-Durán and Rodríguez-Sánchez (2021)stated that the psychological pressure and uncertainty caused by the current changing workplace environment have led to negative consequences for workers. Determine the primary factors that, in the given situation, can lead to employee welfare in terms of employee engagement. A theoretical paradigm to improve involvement during COVID-19 pandemics is put forth through a review of the research. The primary components or the 5Cs are communication, remuneration, cultivation, confidence, and conciliation. Dhanesh and Picherit-Duthler (2021)explains a conceptual framework, the Remote Internal Crisis Communication (RICC) framework, and empirical data to examine the antecedents of employee engagement during remote work in a crisis.

Of all the variables that predicted employee involvement, two-way communication, internal crisis communication objectives and content, and innovative work practices were the ones that predicted it the most. The study also discovered that the association between employee engagement and innovative ways of working was mediated by social connection. Shrimali (2023) states the effects of remote work on organizational dynamics by illuminating the ways in which work arrangements affect employees' levels of engagement. The knowledge gained from this study is important for companies trying to develop a highly motivated and engaged remote workforce while also adjusting to the trends of remote work. Lartey (2022) explains the expansion of existing information by examining the connection between LMX(Leader-Member Exchange), belongingness, and engagement in the context of remote workers. Put differently, the way in which workers view their relationship with their supervisors while they work remotely plays a critical role in maintaining worker engagement. Barhite (2017)states that the strength of the relationship between leaders and members is correlated with employee engagement, according to the study's conclusions, which also showed that perceived organizational communication satisfaction has the strongest association with employee engagement. Shaik and Makhecha (2019) points that the determinants of employee engagement in GVTs (Global Virtual Team) are identified by the study using the Job Demands-Resources theory of employee engagement. Five drivers of employee engagement are conceptualized through interpretive analysis of members' lived experiences





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working in an organization that heavily relies on GVTs to achieve its strategic goals: cultural intelligence, technology, trust, and individual maturity. Kim and Gatling (2018)The study shows that employees intentions to use a hospitality company's VEEP are positively influenced by their perceptions of the tool's use and ease of use. Additionally, the study discovered that employees who were more inclined to utilize their company's VEEP also tended to use it more regularly, which had a favourable impact on their engagement. In due course, the employees who were more engaged shown increased levels of participation and intention to stick around.

NEED FOR EMPLOYEE ENGAGEMENT AND FACTORS AFFECTING REMOTE WORK

Employee engagement can be measured by the degree to which employees feel passionate about their work and how emotionally committed they are to the organization and its mission. The level of employee engagement refers to how the employee perceives his work experience, how that employee is treated in the organization, whether employee feels that his work is suitable. Tenney et al. (2023)The organization is committed to an authentic vision. People have certain needs in their private or professional lives that must be met in order to be engaged, enthusiastic, motivated and committed. For work, some of these needs include technical and managerial skills, autonomy, recognition, a sense of purpose, and a sense of being a valued member of the organization.

Technical competency and Managerial Competency

Technical competencies are the knowledge and skills needed to perform work tasks. They can be learned in an educational setting or on the job and vary by industry. Employers must have the knowledge and skills necessary to do their jobs, and in some cases employers provide these tools or help employees develop them through training or other learning opportunities. Employees must possess general managerial skills in order to advance in their professions. Engaged employees will look for possibilities for professional growth and development that will help them advance in their professions. This is why incorporating continual learning into your organization's culture is critical for maintaining engagement. It helps employees attain managerial competency while also meeting the requirement for growth and development. Employees must possess general managerial skills in order to advance in their professions.

Autonomy and Recognition

Autonomy is a need that businesses must meet from an increasing number of employees, and it is especially vital for remote workers. Autonomy entails enabling, even encouraging, people to work autonomously, providing them with the tools they need, and trusting them to accomplish their jobs successfully. An employee, particularly one with an entrepreneurial drive and an innovative perspective, must be trusted to execute their job properly without micromanagement impeding the process and showing faith and belief in employees' skills, abilities, and decisions. Meeting this need is a key aspect of a strong company culture because it increases job satisfaction, employee engagement, retention, performance, and the quality of work.

Sense of purpose & Feel Valued:

Employee engagement improves when people have a sense of purpose and believe that their work is significant. A shared sense of purpose among coworkers, as well as a connection to an organization's mission, vision, and values, serves as a solid foundation for high levels of engagement. When these conditions are met, firms will witness increased employee engagement, which will result in higher retention rates, enhanced productivity, and increased profitability. Sudden shift from in-person to online work during the pandemic has led to increased work efficiency, online job insecurity and poor adaptation to new ways of working from home. These stressors can drain important social and personal resources, negatively affecting employee engagement.

STRATEGIES To Create A Connection With Virtual Environment ;Impact Of Remote Work On Employee Engagement.

Unambiguous Lines of Communication

Using resources like project management systems, video conferencing, and instant messaging, create clear routes of communication. Promote candid and open communication among team members to foster trust.





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Frequently scheduled video conferences

Plan frequent video conferences to replicate in-person conversations. To keep a personal connection, use video conferences for team meetings, individual check-ins, and even virtual coffee breaks.

Online Team-Building Exercises

Plan social events and online team-building exercises to improve communication. These could be team tasks that foster cooperation and togetherness, trivia evenings, or online games.

Acknowledgment and Gratitude

Reward and acknowledge workers' efforts on a regular basis. Use collaborative platforms, virtual recognition programs, or shout-outs during team meetings to commemorate accomplishments.

Adaptable Work Schedules

Provide flexible work schedules to meet the needs of individuals with varying time zones and preferences. This demonstrates consideration for unique situations and aids in improving work-life balance for staff members.

Employee Welfare Programs

Put into practice employee well-being programs including wellness challenges, mental health courses, and virtual fitness classes. Promote breaks and stress the value of taking time out to refuel.

Opportunities for Professional Development

Offer possibilities for professional growth and virtual training.

Clear Goals and Expectations

Set clear goals and expectations for tasks and projects.

Regularly check in on progress, provide constructive feedback, and ensure that employees understand their roles and responsibilities.

Virtual Mentorship Programs

Establish virtual mentorship programs to facilitate knowledge sharing and career development. Pair experienced employees with those seeking guidance to foster a sense of community.

Technology Enablement

Invest in technology that facilitates collaboration and connectivity.

Ensure that your virtual environment supports seamless communication and access to necessary tools.

Feedback Mechanisms

Implement regular feedback sessions to understand employee concerns and suggestions.

Encourage Informal Interactions

Create spaces for informal conversations and interactions, such as virtual "water cooler" chats or discussion forums.

CONCLUSION

Remote work has revolutionized typical work patterns by allowing individuals to carry out their jobs outside of the usual office setting. After the Covid-19 epidemic, remote work increased in popularity, which was advantageous for many workers. For instance, being able to work from home made it possible for caregivers to balance their personal and professional commitments. Remote employment greatly aided in the promotion of inclusivity and the assistance of people with impairments.. But telecommuting offers support here, as these workers can create an environment





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where they can be comfortable and productive. Financial incentives are not the driving force behind employee engagement in the past, and office perks have lost their appeal in the hybrid and remote work environments that many organizations have moved to. This makes it difficult for managers to find new ways to engage employees in the "new normal". A traditional office environment can be overwhelming for people with social anxiety because they may feel pressured to conform to cultural social etiquette or body language expectations

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Table 1:Age Group

Age	No: of respondents	percentage
18-24	29	24.8%
25-34	46	39.3%
35-44	27	23.1%
45-54	12	10.3%





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55 and above	3	2.6%
Total	117	100%

Table 2:Gender

Gender	No: of respondents	percentage
Male	48	41%
Female	68	58.1%
Prefer not say	1	.9%
Total	117	100%

Table 3:Remote work period

Tuble biltemote Work periou				
Remote work perio	No: of responden	percentag		
Less than one year	33	28.2%		
1-2 years	54	46.2%		
3-5 years	18	15.4%		
Above 5years	12	10.3%		
Total	117	100%		

Table 4:Job Role

Tuble 1.job Note				
Challenges	No of respondents	percentage		
Frontline Employee	34	29.1%		
Middle Management	50	42.7%		
Senior Management	15	12.8%		
Others	18	15.4%		

Table:5

Chi-Square	7.325a
Df	6
P value	0.292

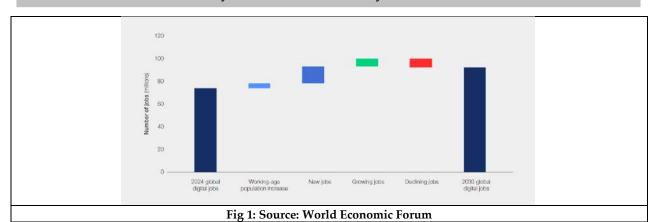
Table 6

Chi-Square	52.785a
Df	12
P value	<.001





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RESEARCH ARTICLE

Green Synthesis, Characterization of Silver Oxide Nanoparticles and its **Photo Catalysis, Biological Studies**

Madhu R¹, Pruthviraj K², Marulasiddeshwra M B³, Sreenivasa S⁴, Sunil K⁵, Suresh D⁶ and Shet Prakash M^{4*}

¹Research Scholar, Department of Chemistry, Tumkur University, Tumakuru, Karnataka, India

²Assisitant Professor, Department of Chemistry, Sri Siddhartha Institute of Technology, SSAHE, Tumkur, Karnataka, India.

³Faculty, Department of Organic Chemistry, Tumkur University, Tumakuru, Karnataka, India.

⁴Professor, Department of Chemistry, Tumkur University, University College Science, Tumakuru, Karnataka, India

⁵Professor, Department of Chemistry, Sri Siddhartha Institute of Technology, SSAHE, Tumkur, Karnataka, India.

6Assisitant Professor, Department of Organic Chemistry, Tumkur University, Tumakuru, Karnataka, India.

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*Address for Correspondence **Shet Prakash**

Professor, Department of Chemistry, Tumkur University, University College Science, Tumakuru, Karnataka, India.

Email: shirsatpm@gmail.com



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ABSTRACT

Ag₂O a versatile nanomaterial having multidimensional application in the field of chemistry with enhanced activity has been synthesized using a endemic regional plant Perioria pinnatum plant having versatile ayurvedic / medicinal activity, the leaf extract has been used as fuel for the green friendly methodology for the synthesis of Ag₂O nanoparticles. Thus synthesized Ag₂O nanoparticles crystalized in FCC system been established through PXRD data and SEM images established the well clear sheet like morphology with mean size of the particles leading to 20nm dimension, followed by TEM images evidenced the spherical shape with smooth edges morphology analysis. Infrared spectrum analysis showed the broad absorption band located around 560 cm-1 corresponds to the due to the Ag-O stretching / lattice vibration. Photocatalytic dye degradation using direct sun light against Methylene blue dye was investigated and it found to be 70% of dye degradation in presence of the nano catalyst, further antioxidant, anticoagulant, antiplatelet and direct hemolytic activities were carried out using well established methods.

Keywords: Ag₂O Nanoparticles, Prioria Pinnatum, PXRD, Photocatalysis, Anti-oxidant, Antiplatelet





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INTRODUCTION

Nanoparticles (NPs) are defined as small particles sized between 1 and 100 nm. Compared with the material bulk state[1].It mainly focuses on the modification of the size, shape and dimensions, has resulted in synthesizing nanoparticles almost equal to Bohr radius [2]. For the synthesis of metal oxides nanoparticles many physical and chemical methods are available in literature. But due to environmental issues and biological applications several authors searching alternative away for the synthesis of metal oxide nanoparticles[3,4]. Recently, introduced an environmentally friendly or bio-based of Green method is one of the simple, inexpensive, non-polluting method and also safe, low-cost, simple protocols when compared to other methods such as spray pyrolysis, sputtering etc. [5,6]. Among this Silver oxide nanoparticles are a very interesting class of metal oxides, silver being a multivalent, it forms various phases like Ag₂O, Ag₀O, Ag₃O₄ and Ag₂O₃ by interacting with oxygen[7]. Experimentally it is found that Ag2O and AgO are the most observable phases. It has also been reported that they decompose at less than 250°C [8,9], and also their wide range of applications in all the fields such as agriculture[10], sensing[11], catalysis[12] and biology activity[13]. Several researchers have reported the green synthesis of metal nanoparticles using leaf extract of Artemisia Herba-Alba[14], Selaginella (Sanjeevini) leaf extract[15], Lawsonia inermis (henna) leaf extract[16], etc. On the basis of literatures survey the present work is to prepare a Ag₂O NPs using of *Perioria pinnatum* plant leaves extract as reducing and capping agent. Because plant extract can reduce metal ions to nanoparticles (NPs) in a singlestep green synthesis process [17,18]. The obtained Ag₂O NPs were cacterized using different techniques, investigate for the decolourization of Methylene Blue (MB) dye under direct sun light irradiation and the results show good photocatalytic performance of these NPs. Further, these Ag2O NPs were used to determine the Antioxidant assay. Anticoagulant, Anti-platelet activity.

Experimental Section & Charazterization Techniques

Collection of Perioriapinnatumleaf and preparation of leaf extract, synthesis of Ag2O -Nanoparticles:

In a brief way, the collected Perioria pinnatum leaves were washed with deionized water, crushed into powered and refluxed at 100° C for $3 \sim 4$ hrs. The obtained plant extract were separated from the filtration process through Whatman filter paper No.42 then filtered (concentrated plant extract) was allowed to cool to room temperature. 2 g of concentrated plant extract was dissolved in 100 ml distilled water and weighed 1.69 g of silver nitrate was added to 30ml of this plant extract solution and heated in a pre-muffle furnace at 400° C, the reaction was completed 4hrs finally to get brown colored Ag₂Onanoparticles.

Characterization of green synthesized Ag₂O Nanoparticles:

Ag2O-NPs were evaluated for the existence of biomolecules by Fourier transform infrared spectroscopy (FTIR, Thermo Fischer Scientific, Waltham, MA, USA). within spectral range of 4000–400 cm1 with a resolution of 4 cm⁻¹. The spectral data were the average of 50 scans over the entire range covered by the instrument. For the analysis, a pellet formed by mixing the dried nanoparticles and KBr was used. The FT-IR spectra of Silver oxide nanoparticles is shown in the **fig.1** The broad absorption band located around 560 cm⁻¹corresponds to the due to the Ag-O stretching / lattice vibration, the bands at 3124cm⁻¹ in the spectra corresponds to O–H stretching vibration indicating the presence of alcohol and phenol from plant extract [19,20]

The nanostructural aspects of the *Perioria pinnatum* leaves silver oxide nanoparticles were characterized by using X-ray diffractometer using equiped Shimadzu 7000 CuK α radiation (λ =1.54 nm). The sample was scanned in the 20 range of 30°-80° with scanning rate of 5°/ min. The XRD pattern for the synthesized Ag₂

O revealed to confirm the four intense peaks at 38.12°, 44.23°, 64.51°, and 77.69° that can be assigned to the plane of {111}, {200}, {220}, and {311}, respectively as in **fig.2** and designated to thethe face-centered-cubic(fcc) system [21,22].





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SEM (Carl Zeiss, USA) fitted with EDS was used to investigate the surface morphology, particle dispersion, and chemical compositions of the prepared silver oxide nanoparticles at a 15 kV accelerating voltage.fig.3 (a,b,c.d,e)depict the SEM images of well dispersed green synthesized Ag₂O NPsat different magnifications the majority of particles were sheet like structure in shape and there were a few oval Ag₂O NPs as well. And the findings clearly represent a mean size of 20 nm.

The size and shape of the silver oxide nanoparticles were obtained by using transmission electron microscopy (JEOL-JEM 2100 LaB6) operated at 200 keV.. **fig.4** (a,b,c.),displays the usual TEM images made using the sample with a 1 mM concentration and various magnifications (50nm, 20nm and 10 nm). The particles were found to have a homogeneous size distribution, an almost spherical shape with smooth edges, and no evidence of aggregation. UV-Vis spectra of synthesized Ag₂O-NPs were recorded in water as solvent after sonication for 10 minutes using Shimadzu UV-1800A11454907691 UV-Visible spectrophotometer, a prominent SPR peak at 430 nm depicts the formation of Ag₂O Nps **fig.5**[23,24].

RESULT AND DISCUSSION

Antioxidant Activity Using DPPH radical scavenging assay

DPPH, a stable free radical with a characteristic absorption at 517-520 nm, was used to study the radical scavenging effects. The decrease in absorption is taken as a measure of the extent of radical scavenging [27]. The activity (RSA) values were expressed as the ratio percentage of sample absorbance decrease and the absorbance of DPPH solution in the absence of extract at 520 nm. The Ag_2O nanoparticles were proved to be inhibiting the DPPH free radical scavenging activity with IC50 value of $55.\mu$ L/mL as show in **fig.6**

Photocatalytic Activity

Silver oxide nanoparticles is very good semi-conductor metal in the field of electrochemical, electronic, optical properties, oxidation catalysis, sensors, fuel cells, photovoltaic cells, all optical switching devices, optical data storage systems. In recent year due to its high catalytic activity, and selectivity, nanoparticles is used as photo catalysts [28]. Therefore in this present work synthesized 20mg of Ag2O is used degrade methylene blue (MB)aqueoussolution(5ppm). The absorbance of standard dye was recorded and the whole setup was kept under sunlight with constant stirring. About 2 ml of suspension was withdrawn ateach 20 min intervals of time and the absorbance was recorded at 665 nm using UV-visible spectrophotometer. The photo catalytic activity methylene blue dye were slowly decrease at specific interval time as show in the fig.7. The degradation efficiency was calculated using the following equation.

% of degradation =
$$\frac{C_i - C_f}{C_i} X 100$$

Where C= Initial concentration of MB, C= Final concentration of MB in the solution afterirradiation for agiven time interval. From the experiment it was found to be for 20mg of Ag₂O degrades 70 % of Methyl blue dye

Direct Haemolytic Activity

Using washed human erythrocytes showed direct hemolytic action. Human packed erythrocytes were combined with phosphate buffered saline (PBS) at a ratio of 1:9 v/v. Following this, dose-dependent additions of 20, 60, and 100 μ g of Ag2O nanoparticles were made to 1 ml of the suspension, which was then kept at 37°C for an hour. Centrifuged at 1000g for 10 minutes at 37°C after the reaction was stopped with 9 mL of cold PBS. The amount of free haemoglobin that was found in the supernatant and measured at 540 nm was used to compute the percentage of RBC cells that were lysed. As shown in **Fig. 8**, Ag2O (0-100 μ g) nanoparticles have a direct hemolytic impact on human RBC blood cells (). As a consequence, it was found that varying concentrations of Ag2O (0–100 μ g) nanoparticles did not hydrolyze on RBC when compared to the positive control (water) and the negative control (PBS) on human erythrocytes [25].





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Anticoagulant Activity

Human platelet-rich plasma (PRP) was prepared using the standard procedure [26]. A platelet concentration of 3.1108 platelets/ml was adjusted for PRP. The material in question was held at 37° C for two hours because of the aggregation process. For all of the aforementioned preparations, plastic or siliconized glassware was utilised. As Quick et al. previously indicated, the plasma recalcification time was calculated. In a 10-mM TrisHCl (20-l) buffer with pH 7.4, 0.2 ml of citrated human plasma was added to the Ag2O (0-100 μ g) nanoparticles and pre-incubated for 1 minute at 37 °C. 20 mL of 0.25 M CaCl2 were added to the pre-incubated mixture after that, and the clotting time was noted. The plasma recalcification time was estimated using human platelet rich plasma, as shown in Fig. 9. (PRP). According to the figure, human platelet rich plasma (PRP) typically clots in 163 seconds. At 20 g, it was discovered that Ag2O's clotting time was 172 s. Clotting time increased to 230s as concentration increased from 20 to 100μ g/mL.

Antiplatelet Activity

Turbid metric approach [27] was combined with the Chronology dual channel whole blood/optical Lumi aggregation system (Model-700) Variable quantities of $Ag2O(50-150\mu g)$ nanoparticles in 0.25ml were used to treat washed platelets. The Aggregation was then started without the use of agonists like ADP, and it was left running for 6 minutes. A significant contributing factor to thrombotic diseases and associated consequences is platelet hyperactivity. Ag2O inhibits the agonist-induced platelet aggregation at a concentration-dependent level in Figure, which was used to evaluate the antiplatelet activity of the compound on platelet function. Agonist-induced platelet aggregation was suppressed by roughly 51%, 73%, and 84%, respectively, at a concentration of 150 μg as shown in Fig.10.

CONCLUSION

In the present work, Ag₂O NPs have been successfully synthesized by solution combustion method using Perioriapinnatum leaf extract because it is easy, economical, non-toxic and eco-friendly method. The synthesized Ag₂O NPs were subjected to wide range of analytical techniques for the confirmation of formation and morphology, further applications shows good photodegradation and antioxidant activity , anticoagulant activity and Direct hemolytic activity.

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Conflict of Interest

The authors declare that there is no conflict of interests regarding the publication of this article.

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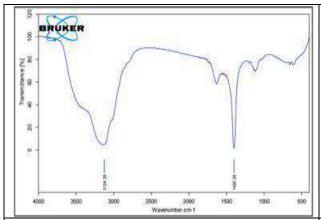
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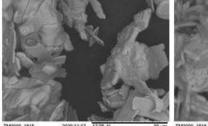
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Fig 1: Infrared spectra of Silver Oxide Nano Particles "Perioria pinnatum" Plant

Fig 2: XRD of Silver Oxide Nano Particles obtained by the addition of Plant Extract.



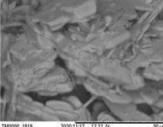
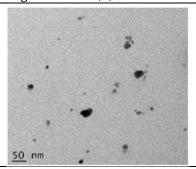
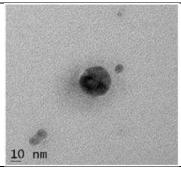




Fig 3: Scanning electron microscopic images of "Perioriapinnatum" leaf extract-silver nanoparticles at different magnifications (a,b,c).





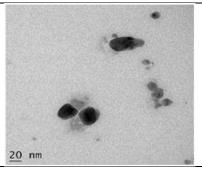


Fig 4: Transmission electron microscope images of "Perioriapinnatum" leaf extract-silver Nanoparticles at different magnifications (a,b,c.).





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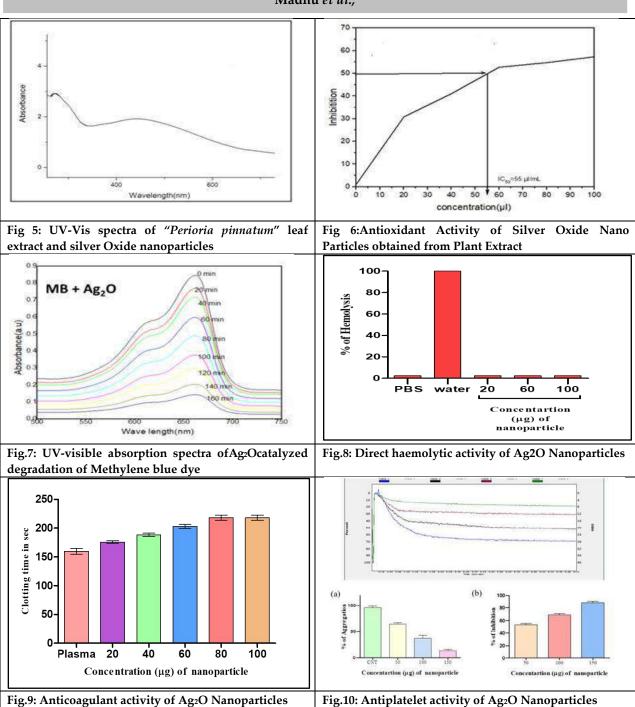




Fig.10: Antiplatelet activity of Ag₂O Nanoparticles



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RESEARCH ARTICLE

Synthesis and Characterization of ZnO Nanostructured Sensors via Co-Precipitation: A Comprehensive Study for Toxic H2S Gas Detection

Vishal S. Kamble¹, Sanjay K. Patil², Shailesh N. Vajekar³, Jyotsna G. Pargaonkar^{1*} and Suresh T. Salunkhe⁴,

¹Assistant Professor, Department of Chemistry, Changu Kana Thakur Arts, Commerce and Science College, New Panvel (Autonomous), (Affiliated to University of Mumbai) Mumbai, Maharashtra, India. ²Principal, Department of Chemistry, Changu Kana Thakur Arts, Commerce and Science College, New Panvel (Autonomous), (Affiliated to University of Mumbai), Mumbai, Maharashtra, India. ³Associate Professor, Department of Chemistry, Changu Kana Thakur Arts, Commerce and Science College, New Panvel (Autonomous), (Affiliated to University of Mumbai) Mumbai, Maharashtra, India. ⁴Principal, Department of Chemistry, Dahiwadi College, (Affiliated to Shivaji University), Maharashtra, India.

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*Address for Correspondence Jyotsna G. Pargaonkar

Assistant Professor,

Department of Chemistry,

Changu Kana Thakur Arts, Commerce and Science College,

New Panvel (Autonomous),

(Affiliated to University of Mumbai)

Mumbai, Maharashtra, India.

Email: jyotsnapargaonkar@gmail.com



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ABSTRACT

In present study, ZnO nanostructures were successfully synthesized utilizing a facile co-precipitation method. A comprehensive analysis was conducted to investigate the structural, morphological, and elemental properties using advanced techniques, including XRD, FESEM, HRTEM, and XPS. Furthermore, the gas sensing capabilities of the ZnO nanostructures were systematically examined against toxic H₂S gas. Remarkably, the ZnO nanostructures exhibited a good response towards H₂S gas, at a concentration of 100 ppm and operating at an elevated temperature of 200°C. The ZnO sensor demonstrated commendable stability in H2S gas detection. The study also proposed a sensing mechanism elucidating the response of prepared material towards H₂S and air. The results collectively present a novel and cost-effective approach for the construction of ZnO nanostructured sensors, offering immense potential for effective H2S gas sensing applications. This work contributes to advancing the understanding of nanostructured metal oxide sensors and their applicability in environmental monitoring systems.





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Keywords: ZnO Nanostructure, HRTEM, H2S gas, Sensor.

INTRODUCTION

Gas sensing capabilities of semiconducting metal oxides, such as ZnO, Fe₂O₃, V₂O₅, SnO₂, WO₃, In₂O₃, TiO₂, and Ga₂O₃ make them auspicious materials for detecting various toxic and pollutant gases [1-2]. ZnO, in particular, featuring a direct bandgap of 3.37 eV, high electron mobility (210 cm²V⁻¹S¹), and a substantial excitonic binding energy (60 meV). ZnO's structural diversity is another advantageous feature. Its versatile crystalline structures and morphologies enable the customization of sensor designs, allowing for the optimization of performance based on specific gas detection requirements. This adaptability contributes to the material's effectiveness in detecting a wide range of gases [3-6]. As a chemo-resistive nanostructure, ZnO has proven effective in detecting various gases, including H2S, NO2, H2, CO, NH3, CH4, LPG, ethanol, and acetone. Toxic gases, when released into the environment, pose significant threats to human health. Inhalation of these noxious substances can lead to respiratory problems, irritation of the eyes and skin, and, in severe cases, can result in long-term health issues or even fatalities. Consequently, the ability to promptly recognize and respond to the presence of toxic gases is paramount for minimizing the risks to individuals and communities [7]. Indeed, among various pollutants, hydrogen sulfide (H2S) emerges as a prominent and highly toxic air pollutant. This compound takes center stage due to its widespread occurrence, often generated through combustion processes in industries [8]. Inhaling a low concentration of H₂S gas can induce symptoms such as dizziness, nausea, and headaches in humans, while exposure to higher concentrations exceeding 250 ppm can lead to severe illness and even fatalities. An estimated three million tons of H2S gas are discharged into the environment annually. Although the characteristic rotten egg odor becomes noticeable to humans at concentrations above 250 ppm, detection becomes challenging below 100 ppm. Consequently, it is imperative to develop methods for H₂S gas detection below the 100-ppm threshold to ensure both human safety and environmental protection[9-10]. Among various synthetic methods for creating ZnO nanostructures, the coprecipitation is a simple technique. The pressing demand for highly sensitive, selective, and stable gas sensors capable of detecting H2S gas at deficient concentrations with rapid response and recovery times is evident.

Various synthetic techniques have been explored to produce ZnO nanostructures, each with its advantages and drawbacks. Notably, the co-precipitation method proves to be particularly advantageous due to its ease of implementation, cost-effectiveness, and efficiency in yielding nanostructures with desirable properties. In the scope of this research endeavor, our primary focus is directed towards the synthesis of ZnO nanomaterials utilizing the coprecipitation method. The specific objective driving this synthesis is the development of a highly sensitive H₂S sensor designed to detect lower concentrations within the environment. To attain the goal, a comprehensive characterization process was implemented, using a various array of analytical techniques, including XRD, EDAX, FESEM, HRTEM, and XPS. These techniques were instrumental in authentically validating the surface and morphology of the created ZnO nanorods. Our investigation extends beyond mere synthesis and characterization, delving into the intricate realm of the ZnO nanorodsH2S sensing capabilities at varying concentrations and temperatures. This comprehensive study permits for a nuanced understanding of the sensor's performance under diverse conditions. Furthermore, we meticulously examined various critical sensing parameters, including stability, reproducibility, gas response, and recovery time. Through this systematic analysis, the obtained results unequivocally establish the ZnO nanosensor's exceptional potential, particularly when operated 100 ppm H2S concentration and 200°C temperature. In essence, our research underscores the significant strides made in the development of a ZnO nanorod-based H2S sensor, presenting compelling evidence of its promising performance characteristics. The implications of this study resonate in the realm of high-performance gas sensing technologies, with the ZnO nanosensor poised to make substantial contributions to H2S detection in the future.





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MATERIALS AND SYNTHETIC METHOD

Zn(CH₂COO)₂·2H₂Oand m-cresol AR grade and 80% ethanol were gained commercially. The preparation of ZnO nanomaterial involves a meticulous procedure, beginning with precursor preparation through the dissolution of zinc acetate. The subsequent stages include nucleation, annealing, and powder crushing to attain a fine ZnO powder. The formation of a sticky gel in m-cresol follows, with a detailed stirring process. The final step entails the deposition of this gel onto a glass substrate through the drop-casting method, thereby forming the foundation for the fabrication of a thin film featuring ZnO nanostructures. A schematic representation (Fig. 1) encapsulates the entirety of this intricate synthesis scheme.

Characterization and Gas sensing tests

XRD analysis was conducted utilizing a Bruker D2 Phaser with $CuK\alpha$ radiation (wavelength of 1.54056 Å). The analysis spanned a 20 range of 20–80°. For an in-depth exploration of the surface morphology and particle size, FESEM, and HRTEM analyses were employed. Additionally, XPS was employed to investigate the atomic composition of the ZnO sample. The capabilities of gas sensing of the ZnO material were thoroughly examined using a custom-designed gas sensor unit, incorporating the Keithley 6514 electrometer for accurate data recording. The choice of M/s Shreya Enterprises as the gas supplier ensures the credibility of the experimental setup, contributing to the validity of the findings in assessing the ZnO-based sensor's performance across various target gases. The sensitivity (S) of the sensors calculated through a specific relation.

$$S(\%) = \frac{[Ra - Rg]}{Rg} \times 100 \tag{1}$$

where, Rg and Ra are resistance the target gasand the air, respectively.

RESULT AND DISCUSSION

The X-ray diffraction (XRD) technique serves as a pivotal method for conducting structural analysis. Fig. 2 showcases the XRD pattern of ZnO nanomaterials. The distinct peaks at 31.98°, 34.63°, and 36.47°, are identified with corresponding crystallographic planes, namely (100), (002), and (101) planes respectively. These findings indicate a hexagonal wurtzite structure for the ZnO nanorods, a conclusion supported by the coordination with JCPDS card(01-079-0205) (Inset of Fig. 2.)The meticulous investigation into the structural aspects of ZnO, coupled with the absence of impurity peaks and the utilization of the Scherrer relation for crystallite size determination, collectively enhances the depth and precision of the structural characterization[11]:

$$D = \frac{\hat{K} \lambda}{\beta \cos \theta}$$
 (2)

Furthermore, lattice constants a and c as well as dislocation density (δ) were calculated using specific equations, the details of which are not provided here, and the results were tabulated in Table 1 [12-14].

$$\frac{1}{d^2} = \frac{4}{3} \left(\frac{h^2 + k^2 + hk}{a^2} \right) + \frac{l^2}{c^2} \tag{3}$$

Dislocation density (
$$\delta$$
) = $\frac{1}{D^2}$ (4)

This comprehensive analysis not only validates the crystalline nature and purity of the ZnO nanorods but also provides valuable insights into their structural dimensions and parameters through precise calculations based on the XRD data.

Morphological studies

FESEM and HRTEM analysis

In Fig. 3(a), the top-down view of FESEM images of 100 nm ZnO illustrates stepped nanorods featuring hexagonal nanostructures that are distributed in a random manner. This alignment is in agreement with the XRD results. The average length of ZnO NRs is found to be in 120-380 nm, while their diameter measures approximately 40-110 nm. This particular morphology contributes significantly to the exceptional gas response observed in ZnO NRs when





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applied in gas sensing applications. To gain further morphological study of ZnO NR's, HRTEM was employed. The typical HRTEM copy of the 50 nm ZnO NR's are obtainable in Fig. 3(b), revealing stepped nanorods with a well-defined crystalline nature, consistent with the findings from FESEM images. Furthermore, HRTEM analysis emphasizes the interparticle permeability, underscoring the porous nature of ZnO nanorods (NRs). This porous structure and additional surface area are a noteworthy characteristic that significantly enhances sensing capabilities of ZnO NRs by providing additional surface area available for adsorption.

XPS analysis

In Fig. 4a, the XPS survey spectrum reveals solely Zn, O, and C peaks, underscoring the sample's high purity. Moving to Fig. 4b, the Zn 2p scan shows two distinct peaks at 1044.41 and 1021.26 eV, corresponding to Zn2p1/2 and 2p3/2 respectively. The observed energy difference of 23.15 eV aligns of the Zn^{2+} valence state, corroborating well with established reference standards [15-16].

GAS SENSING PERFORMANCE:

Gas response

The gas sensing capabilities of a pristine ZnO nanostructure were evaluated across a range of H₂S concentrations (5 ppm to 100 ppm) with the temperature of 200°C. The ZnO sensor exhibited a remarkable 164% enhancement in gas response when exposed to a concentration of 100 ppm. Notably, the response prepared sensor to H₂S displayed a linear correlation with concentration, escalating from 25 to 100 ppm. At the highest tested concentration of 100 ppm, the ZnO demonstrated a superior response.

Response/recovery time

In Fig. 5, the variations in response/recovery times of the ZnO sensor in relation to H₂S gas concentration (25, 50, and 100 ppm) at temperature of 200°C are illustrated. Notably, the response time exhibits a decrease from 32 to 15 seconds, whereas the recovery time demonstrates an increase from 61 to 114 seconds with the rising concentration of H₂S gas. This trend indicates a correlation, either inversely or directly, between the response/recovery time and the concentration of H₂S gas. The observed decrease in response time can be attributed to the higher concentration of H₂S gas facilitating a more rapid interaction with the sensing material. As the gas concentration increases, a greater number of H₂S molecules become accessible to the sensing material, leading to a quicker response time. This phenomenon is indicative of the sensor's heightened reactivity to higher concentrations of H₂S. Conversely, the increase in recovery time is associated with the extended duration required for the complete desorption of H₂S gas from the sensing material. Higher gas concentrations may lead to a larger quantity of H₂S molecules adsorbed onto the sensor's surface, necessitating a longer time for desorption and restoration of the baseline signal. This finding underscores the importance of considering both response and recovery times for a comprehensive evaluation of the sensor's performance.

Stability

In Fig. 6, the graph illustrates the durability of the prepared ZnO sensor. The study focused on evaluating the stability of the zinc oxide sensor in response to H₂S gas over 50 days at the temperature of 200°C, with measurements conducted every 10 days for 100 ppm concentration. Notably, the ZnO sensor exhibited 52% stability. It is remarkable that the initial sensor response decreased from 164% to 86% during the 50-day duration but eventually stabilized. This stabilization is attributed to the impact of humidity on the sensor. These results emphasize the promising applicability of ZnO as a material for gas sensors in the context of detecting H₂S gas.

Sensing mechanism

Figure 7 illustrates a schematic representation of the potential gas sensing mechanism employed by the ZnO sensor. In the presence of the air, oxygen absorbed onto the ZnO sensor surface. These oxygen molecules undergo conversion into various oxidation states (O-, O2-, and O2-). This process results in the formation of a free-electron region (electron depletion layer). Consequently, there is a simultaneous decreases number of electrons and an increase in the barrier height for electron transport, leading to a high resistance state for the ZnO sensor (Fig. 7a)





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[17].It's worth noting that O₂ ions are stable under 100°C, O between 100°C and 300°C, and after 300°C, O² ions become dominant and integrate into the lattice [18-19]. Fig. 7b illustrates the introduction of H₂S gas to the prepared sensor. H₂S serves as a reducing gas, and when samples are exposed to H₂S at elevated temperatures, it facilitates the removal of adsorbed oxygen. This process effectively restores electrons to the conduction band, resulting in a reduction in the resistance. This phenomenon crucially hinges on the interaction between H₂S and oxygen ions (O⁻) at the surface of ZnO. Specifically, the interaction re-injects electrons into the conduction band, leading to an enhancement in conductivity compared to the baseline state (20-21). Therefore, exposure to H₂S at higher temperatures induces a beneficial effect on the electrical properties of the semiconductor by promoting the removal of adsorbed oxygen and facilitating electron flow in the conduction band.

CONCLUSION

In this study, a co-precipitation technique was employed to successfully synthesize ZnO nanostructures. XRD analysis confirmed the hexagonal wurtzite structure. The FESEM study revealed a porous and uniform morphology in pure ZnO, making it well-suited for gas sensing uses. Based on comprehensive gas sensing property assessments, it was determined that the ZnO nanomaterial exhibited a remarkable maximum response of 164% at 100 ppm H₂S gas and at of 200°Ctemperature. The prepared sensor demonstrated favorable response/recovery times, and stability Consequently, ZnO nanostructures emerge as highly promising materials for upcoming sensors designed for detecting H₂S gas at a concentration of 100 ppm, showcasing exceptional capabilities in the realm of gas sensing technology.

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CONFLICT OF INTEREST

The authors assert that they have no discernible conflicting financial interests or personal affiliations that could be perceived as influencing the research presented in this paper.

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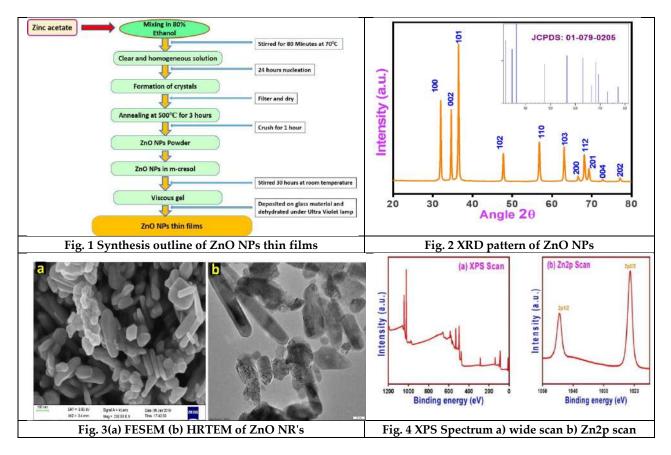


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Table 1: Structural parameter of prepared ZnO sample

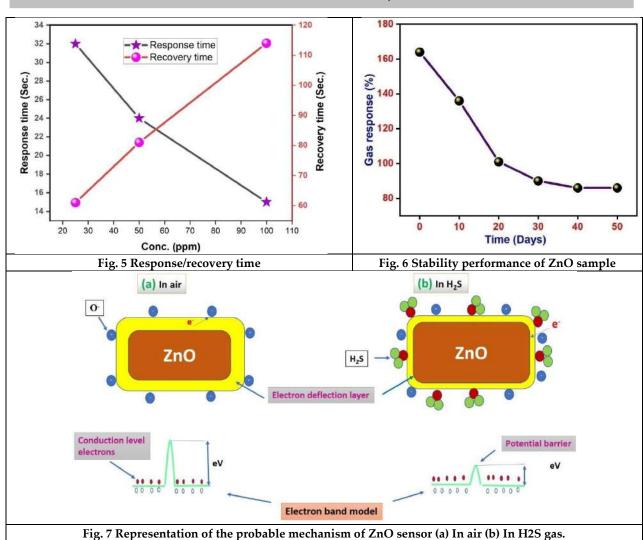
Material D a c ZnO NPs 34.7 nm 3.244 5.182







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RESEARCH ARTICLE

Nanoparticles as Food and Medicine: a Systematic Analysis of Recent Nanotechnology Developments in Fishery Industry

Vasantharaja D*, Nagaraj S, Peetha Sarika and Vimala V

Department of Zoology, Tagore Government Arts and Science College (Affiliated to Pondicherry University) Puducherry, India.

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*Address for Correspondence

Vasantharaja D

Department of Zoology,

Tagore Government Arts and Science College (Affiliated to Pondicherry University)

Puducherry, India.

Email: vasaini20@gmail.com



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ABSTRACT

The unique properties of nanoparticles (NPs) have driven the rapid growth of nanotechnology as a field of study. In both human and animal healthcare, NPs find applications in diagnostics, vaccinations, drug delivery, and gene therapy. This systematic review investigates the cutting-edge advancements of nanotechnology in the fisheries industry, focusing on its diverse applications in food and medicine. NPs are emerging as transformative tools, offering distinct advantages in improving food production, preservation, and drug delivery for aquatic resources. This review provides an overview of NPs utilization, with a particular emphasis on their antimicrobial properties and their role as supplementary feed in the fishing sector. Research in fish vaccines and medications is extending to include NPs for combating various bacterial, fungal, and viral infections. The aquaculture industry is increasingly exploring NPs as a targeted food source and reliable diagnostic tool for various ailments. Nevertheless, it is essential to acknowledge that there are numerous untapped nanotechnologies in the fisheries sector. This study underscores the vast potential of NPs in revolutionizing the fishery industry while highlighting the need for further exploration and innovation in this promising field.

Keywords: Nanotechnology, Nanomedicine, Nanoparticles, Fish medicine, Aquaculture

INTRODUCTION

The term "nanotechnology" refers to the field of study dedicated to the construction and manipulation of subatomic particles. In recent years, tools have been developed for investigating and controlling materials at the nanoscale, typically ranging between approximately 1-100 nm. The distinctive properties of these particles open up avenues for innovative applications [1]. Artificial nanoparticles (NPs) are deliberately synthesized, while naturally occurring





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ultrafine nano-sized particles represent one category of nanomaterials [2]. Nanomedicine, encompassing both human and veterinary healthcare, involves the application of nanotechnology. It focuses on the research and development of NPs and nanodevices for medical applications [3, 4], including their use in fish feed. Nanoparticles are available in various forms, each finding medical applications as outlined in Figure 1 [4]. Nanospheres, characterized by their small size and spherical shape measuring only a few nanometers, are particularly advantageous for drug delivery due to their extensive surface area [5]. They also demonstrate applicability in tissue regeneration [6]. Another type of nanoparticles, nanocapsules, possesses a nanoscale shell encapsulating a core, safeguarding the enclosed medicine whether oil or water from degradation and hydrolysis [7]. Liposomes, nano-sized lipid bilayer spheres resembling eukaryotic cell membranes, are well-suited for transporting both lipophilic and hydrophilic drugs [3, 8]. Carbon nanotubes, which can be single or multiple-walled, are another type of tiny particles used a lot in medicine. They have a lot of space on their surface and can get into cells easily, like tiny needles. This makes them great for carrying drugs, including ones that fight tumors [9]. However, studies show they can cause blood clots in blood vessels [10]. Dendrimers are another kind of tiny particles. They're like tiny three-dimensional structures with branches coming out from a center. They are light and have many branches on their surface, which makes them useful for growing tissues, fighting germs, and carrying genes, vaccines, and medicines [11, 12]. Nanoparticles (NPs) are increasingly being utilized in fisheries as medicine, and their recent application extends to diverse areas, such as integrating them in aquaculture as feed. These tiny particles serve various functions, including acting against microbes, transporting drugs or genes, delivering vaccines, and accurately diagnosing fish infections [13]. Compared to bulk materials, nanoparticles offer numerous technical and biological advancements. They have special qualities due to their tiny dimensions and massive particular area of surface. However, these fields of nanotechnology research are still relatively underdeveloped. In this review article, we present a thorough overview of both proven and potential uses of nanomaterials in the fish feed and fish medicine industries.

MAJOR SYNTHESIS OF NANOPARTICLES

The production of nanoparticles can be achieved through two primary methods: the descending or ground-up approach [14]. In the top-down technique, conversion of large-scale and micro-scale metal into nanoscale particles is accomplished through mechanical grinding. Following this, calming substances like colloidal defending substances are introduced to prevent the NPs from oxidizing or reassembling at the micro-scale [15]. Various chemical and physical processes, including electro-chemical reduction of metals and sono decomposition [16], can be employed to fabricate nanoparticles, which are categorized as bottom-up methods.

Physical and Chemical synthesis of NPs

Thermal combustion of organic solvents can be used for NP production [17]. Cryochemical synthesis may produce metal NPs with a diameter ranging from 5 nm to 80 nm [18]. To create silver nanoparticles (Ag-NPs), scientists have turned to microwaves, specifically for the physical breakdown of Ag at distinct wavelengths of microwave rays [19]. In comparison to the thermal approach, this technique was faster and produced an elevated attention of Ag-NPs under the same conditions. To generate a larger particle size, Jiang et al. [19] observed that the reaction time and temperature needed to produce the silver nitrate were also increased, along with the quantity of silver nitrate employed. Silver particles ranging between 15 and 25 nm in size were produced by using high concentrations of polyvinyl pyrrolidone (PVP). A number of methodologies are utilized for the making of NPs. These embrace sustained ablation with laser [20], electro-reduction of silver nitrate in polyethylene glycol in an aqueous solution [21], ember ejection [22], micro-emulsion for creating silver- Iron (II,III) oxide-NPs [23], and the AgNO3 undergoes chemical reduction with trisodium citrate in the presence of sodium tetrahydridoborate (III) as a reluctant to synthesize PVP-coated Ag-NPs [24, 25]. Additionally, methods for synthesizing nonmetallic NPs, like polymeric chitosan (CS)-NPs, have been demonstrated through various approaches. Ionotropic gelation ia a techniue in which, chitosan (CS) is dissolved in acetic acid, followed by the addition of the polyanion tripolyphosphate (TPP). This process relies on the magnetized connection among the positively charged amine group of CS and the negatively charged polyanion polymer [26, 27]. In the polyelectrolyte cluster technique, the magnetized connection between the positively charged groups in the CS and the DNA is crucial, leading to charge neutralization during NPs production





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[26]. The micro-emulsion method [28], which utilizes surfactants, has drawbacks such as the utilization of natural substances, an extended manufacturing duration, and the intricate process of cleaning operations. Precipitation [29] and double-emulsion solvent evaporation [30, 31] are typically the two steps in the synthesis of PLGA-NPs. In the process, polyelectrolytes with positive and negative charges are mixed together, leading to the formation of a complex structure held together by electrostatic forces [26]. This technique is called polyelectrolyte complex allows for precise control over the properties of the resulting material, making it suitable for a wide range of applications. The polyelectrolyte complex technique involves the formation of a stable material through the interaction of oppositely charged polymers. This technique is used in various applications such as drug delivery, tissue engineering, and surface modification. The micro emulsion technique utilizes surfactants and also has many drawbacks [2]. Production of PLGA by precipitation [29] followed by double emulsion evaporation [30, 31]

Biological / Green synthesis of NPs

Researchers are very interested in finding ways to make NPs that are good for the environment and don't cost a lot of money. The use of biological methods is seen as a key part of this strategy's success [32]. Bacteria, fungus, and plants are the primary sources of biologically produced NPs. Bottom-up reduction and oxidation reactions are important to the biosynthesis of NPs [33]. Microorganisms or plant chemicals that have antioxidant or alleviating objects catalyze predecessor molecules to make the preferred NPs. A biosynthetic nanoparticle system typically includes three main components: This substance serves as a solvent in the process of synthesis, functions as an ecologically conscious reducing agent, and acts as a secure stabilizing agent [33]. Ag-NPs were made using liquid extracted from Origanum vulgare leaves. These particles showed the ability to kill bacteria and were toxic to lung cancer cells [34]. Both silver (Ag) and gold (Au) nanoparticles were created using liquid extracted from cashew nut shells. These particles can kill bacteria that cause fish infections [35]. Additionally, high amounts of Ag-NPs made with Camellia sinensis a tea leaf extract were able to kill Vibrio harveyi bacteria in young Fenero penaeus indicus shrimp [36]. A recent study looked at silver nanoparticles (Ag-NPs) made from Origanum vulgare leaves and compared them with chemically made ones. The two kinds were examined utilizing UV-spectrophotometry, transmission electron microscopy (TEM), and dynamic light scattering. The bacteria Streptococcus agalactiae, Aeromonas hydrophila, and Vibrio alginolyticus, as well as the fungus Aspergillus flavus, Fusarium moniliforme, and Candida albicans, were tested in various fish species [37]. Additionally, zinc oxide (ZnO) nanoparticles were made in a broth from Aloe leaves and showed stronger germkilling ability compared to regular ones [38]. This eco-friendly and cost-effective method involved making ZnO-NPs with the help of Aeromonas hydrophila bacteria, resulting in nanoparticles that can kill bacteria and fungi [39].

ANTIBIOTIC-RESISTANT BACTERIA

Antibiotics have proven effective in combating bacterial infections for an extended period. Nevertheless, the uncontrolled and excessive utilization of antibiotics may contribute to the rise of microorganisms that are resistant to antibiotics [40-42]. These bacteria no longer respond to the effects of antibiotics. The transfer of resistance genes between terrestrial and aquatic bacteria is feasible due to shared components in their mobile genomes. Such gene transfer poses potential risks to the health of both animals and humans [42-44]. Tuevljak et al. conducted a survey across 25 countries to assess the antimicrobial usages in fish farms [45]. Their findings revealed that the most commonly employed antibacterial agent is tertracycline. The excessive application of antibiotics in aquaculture is significantly correlated with proliferation of antibiotic-resistant bacteria. This increase also involves a rise in the number of bacteria that are resistant to different treatments, like methicillin-resistant *Staphylococcus aureus* (MRSA), as well as some microorganisms that are opposing to manifold and extremely powerful drugs [45, 47]. Certain *Aeromonas hydrophila* found in farmed tilapia were resistant to a wide range of antibiotics [48]. Additionally, antibiotic resistance was seen in other bacteria stains [43, 49]. Di-Cesare and colleagues found antibiotic-resistant enterococci in the sediment of fish farms, raising concerns about the spread of resistance to strains that can affect humans [50, 51].





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INNOVATIVE ANTIBACTERIAL APPLICATIONS OF NANOPARTICLES IN FISH MEDICINE

The emergence of drug-resistant microbes in aquaculture has become an increasingly concerning issue. Consequently, scientists have explored the possibility of utilizing NPs as a potential antibacterial solution (Table. 1) [37, 49]. Metal NPs have potent antibacterial effects against many microorganisms [52]. Following this, we will discuss research into the use of metal NPs in combating fish diseases

Ag-NPs

Nano-antibacterial agents have been studied extensively, but Ag-NPs have received the most attention. In contrast to antibiotics, which typically only operate through one mechanism [53–55], multi-target antimicrobials are able to circumvent bacterial resistance by targeting numerous pathways at once. A method involving the silver ions (Ag+) released has been identified [53]: Silver ions interact with proteins present in the bacterial cell membrane, leading to membrane rupture and eventual cell death [58]. When Ag+ enters a cell, it attaches to mitochondria and nucleic acids, causing harm and interrupting replication of cells [59]. Ag-NPs demonstrate potent antibacterial activity, particularly against MDR bacteria, as highlighted by Prakash et al. [56]. Notably, methicillin-resistant *Staphylococcus aureus* (MRSA) has been found to be susceptible to the bactericidal effects of Ag-NPs [56]. When Ag-NPs are created using lemon juice as a dropping agent, they show effectiveness against *Staphylococcus aureus* and *Edwardsiella tarda* bacteria. Additionally, they exhibit anti-cyanobacterial effects against Anabaena and Oscillatoria species [49]. The studies have utilized the extract of leaf buds from the mangrove *Rhizophora mucronata* for the green synthesis of Ag-NPs, aiming to demonstrate antibacterial activities [60].

These eco-friendly or "green" Ag-NPs exhibited effectiveness comparable to conventional antibiotics but with significantly fewer negative side effects. Notably, long-term therapy with Ag-NPs resulted in a 71% reduction in mortalities in *Vibrio harveyi*-infected *Feneropenaeus indicus* at increased doses of Ag-NPs [36]. Ag-NPs showed potent antifungal activity against *Candida* species, with antifungal effects comparable to those of the commercial antifungal Amphotericin B [61, 62]. Researchers have found that Ag-NPs have antifungal efficacy against dermatophytes [63]. In another study, Ghetas et al. demonstrated that the Ag-NPs inhibit the growth of three fungal strains isolated from Nile tilapia) tested in their study. The results indicate that the Ag-NPs synthesized through biological means have shown better efficacy than those synthesized chemically. Furthermore, it is promising to note that the antimicrobial effectiveness of Ag-NPs against the fungal strains can be improved with an increase in the administered dose [37]. Ag-NPs can bind to HIV-1 virus proteins *in vitro* [64], demonstrating their anti-viral capabilities. The influenza A virus can be inhibited by either Ag-NPs or Ag-NPs -CS composite [65, 66]. When it comes to Ag-NPs antifungal and antiviral properties in fish medicine, there is a dearth of published work.

Au-NPs

Due to the non-hazardous nature of Au-NPs to eukaryotic cells, considerable research efforts are being directed towards studying their antimicrobial activities [67]. Au-NPs exhibit the ability to act together with biological macromolecules as well as non-proteins and they can perform various functions in biology [68]. The bactericidal effects of gold nanoparticles supported on aluminium silicate were demonstrated against *E-coli* and *S. typhi* [69]. Functionalized gold nanoparticle was established to suppress the growth of multidrug-resistant bacterial strains [67]. Additionally, antibacterial activity was observed in Au-NPs produced through "green" synthesis against a bacterial strain isolated from fish [35]. Au-NPs exhibit antibacterial properties, and they can employ three different mechanisms to kill microbes. The first mechanism involves altering the cell membrane of specific microbes, disrupting the oxidative phosphorylation process. This alteration result in reduced efficiency of the F-type ATP synthase, leading to a decrease in the overall rate of ATP synthesis and metabolism. The second mechanism involves modifying how t-RNA attaches to the two subunits of the ribosome. The third strategy involves enhancing chemotaxis [70]. Research has established that Au-NPs can exert fungicidal effects against various Candida species. Interestingly, the antifungal activity of Au-NPs is more pronounced when the particles are smaller. Their effectiveness has been demonstrated to be directly proportional to their particle size [71, 72].





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ZnO-NPs

ZnO-NPs have garnered significant attention for their ability to effectively eliminate bacteria and fungi [38, 49]. These nanoparticles induce damage to the microbe's cell membrane, ensuing in the leakage of cytoplasmic contents from the cell [73]. In the area of fish healthcare, Zn-NPs have demonstrated the capability to restrain the development of various bacterial strains [49]. In a study by Ramamoorthy et al. [74], they looked at how zinc oxide nanoparticles can kill harmful *Vibrio harveyi* bacteria. They found that these nanoparticles were better at killing bacteria compared to regular zinc oxide. Another interesting study involved making ZnO-NPs using a bacterium called *Aeromonas hydrophila*. These nanoparticles were able to kill not only *Vibrio harveyi* but also other microorganisms like bacteria and fungi [39].

Titanium dioxide (TiO2) NPs

TiO₂ NPs infused with magnetic Fe₃O₄-nanoparticles exhibited antibacterial activity, when exposed to light [75, 76]. Due to the affinity of fish pathogens for these nanoparticles, allowing for their easy removal from water using a magnet, these particles can be utilized in water disinfection processes. Additionally, Jovanovic et al. [77, 78] reported that Titanium dioxide nanoparticles have an impact on the fish defence system by suppressing the bactericidal response of fish neutrophils. This makes fish highly vulnerable to disease, leading to increased mortality rates, principally during spreading of diseases, posing a significant concern.

THE USE OF NANOPARTICLES AS CARRIERS FOR THE DELIVERY OF DRUGS AND GENES

A good drug delivery system should have certain important qualities. These include being safe, compatible with the body, able to break down naturally, and keeping the drug stable. It should also target specific areas and have few or no side effects (Fig.2) [4]. Nanoparticles (NPs) have become popular for this purpose because they are very small and can move through barriers in the body like the BBB. Also, NPs have a large surface area compared to their size, which makes them very reactive with different substances [4, 79, 80]. In fish medicine, Chitosan NPs and PLGA-NPs are being studied a lot as carriers for delivering drugs.

Chitosan NPs for drug delivery

Chitosan NPs stand out as favorable choices for drug delivery vehicles due to their exceptional characteristics. Comprising a polymer that is biocompatible, non-toxic, and biodegradable, these nanoparticles are easily eliminated by the kidneys [4]. Their muco-adhesive efficiency makes them well-suited for controlled and sustained drug release [80]. In a study involving *Oncorhynchus mykiss*, for instance, chitosan NPs were used to enhance the delivery system for vitamin C. The synergistic association between chitosan and ascorbic acid resulted in sustained vitamin release for up to 48 hours after exposure, stimulating the fish's inborn immune system [82]. In a different test, hormones in *Cyprinus carpio* (common carp) were given using chitosan nanoparticles (NPs). One group got LH-RH mixed with chitosan-coated gold nanoparticles (Au-NPs), and we compared the results with a group getting just one hormone injection. Both groups had higher levels of hormones in their blood, and the hormones were released more slowly compared to the group with only one injection. Interestingly, injections of hormone mixed with chitosan NPs and hormone mixed with chitosan-coated gold NPs increased the rate of fertilization of eggs by 87% and 83%, respectively, compared to giving repeated injections of luteinizing hormone-releasing hormone alone, which only resulted in a rate of fertilization is about 74% [83].

PLGA-NPs for drug delivery

Polylactic acid and polyglycolic acid are combined to form PLGA, a copolymer. It is non-toxic, biodegradable, and biocompatible. The FDA has given its approval. As a result, numerous researchers have looked at whether using PLGA as a drug carter is feasible [84, 85]. In a recent study, Poly (lactic-co-glycolic acid)-(PLGA)-NPs were loaded with the anti-bacterial medication rifampicin and then injected into zebra fish embryos. Zebra fish embryos are apparent, making it possible to use non-invasive imaging to assess the treatment's effects on Mycobacterium marinum-infected cells. In comparison to rifampicin alone, the rifampicin-PLGA-NPs had a better restorative efficacy towards M. marinum and advanced embryonic survival [86].





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Fish vaccines based on nanotechnology

A fascinating area of medical research is the incorporation of NPs in vaccination compositions. Because of their many benefits as vaccine delivery systems, polymeric NPs have received the greatest attention [87, 88]. These benefits include the capacity to assure antigen stability against enzyme degradation, maintain immunogenicity, and provide prolonged release of the vaccine. Both immune-stimulant adjuvants and targeted antigen delivery nanovaccines have the ability to slowly release antigens [88]. Various types of nanoparticles (NPs) have been used to deliver vaccines, including virus-like organisms, liposomes, immunostimulant complexes, metal NPs, and polymeric NPs [88, 89]. Challenges in making stable nano-vaccines, worries about possible harm, and not knowing enough about how they spread in the body have been the main concerns [89]. In the world of fish vaccine development, polymeric chitosan and PLGA-NPs have been the focus of most research so far. In the inactivated virus vaccine for infectious salmon anemia virus (ISAV), the vaccine contains the DNA code for the ISAV replicas, and chitosan NPs are added to enhance the immune response. This vaccine has been shown to be highly effective, providing up to 77% protection against ISAV [90]. Another vaccine was developed to protect Asian sea bass from the bacteria *Vibrio anguillarum*.

This vaccine contains DNA mixed with chitosan and chitosan/tripolyphosphate nanoparticles. However, this nanovaccine only provided a moderate level of defense against the microorganisms [27, 91]. An oral DNA vaccine was formulated by loading the outer membrane protein K gene of *Vibrio parahemolyticus* onto chitosan NPs. This approach facilitated the delivery of the gene as a vaccine. The resulting recombinant nanovaccine demonstrated the ability to induce a protective immunological response against *Vibrio parahemolyticus* in black seabream (*Acanthopagrus schlegelii*) [92]. Researchers looked at how well r-DNA-chitosan NPs could defend shrimp from the White Spot syndrome Virus (WSSV). It was discovered that the vaccination improved shrimp defense system and generated a defending activity against WSSV when it was administered in the form of an oral supervision [93, 94]. PLGA-NPs have demonstrated effectiveness both as carriers for DNA vaccines and as adjuvants in medicine [30, 95, 96]. However, there has been limited success in utilizing PLGA-NPs in fisheries. For instance, a DNA vaccine for protecting Japanese flounder towards lymphocystis disease virus (LCDV) was formulated, and it could be administered orally [97]. In another study, rainbow trout were victimized against the contagious hematopoietic necrosis virus using an orally delivered DNA vaccine, resulting in a successful elicitation of an immunological response in the fish [31].

THE USE OF NANOPARTICLES IN DETERMINING THE PRESENCE OF FISH PATHOGENS

The use of NPs in diagnostic testing has been given the general term "nanodiagnostic" [98], which is also the name of a specific category of diagnostic procedures. These methods make it possible to make accurate diagnoses of diseases in a short amount of time. One of the NPs that is used in the industry the most commonly is Au-NPs, which may be utilized in a range of diagnostic ways [99, 100]. Au-NPs are also one of the most expensive NPs.

Diagnostics for Fish Bacterial and Fungal Diseases

To specifically diagnose furunculosis in fish tissues, the first use of Au-NPs for pathogen detection was using an A. salmonicida antibody and Au-NPs coupled [101]. For the purpose of detecting fish diseases, this was the initial application of Au-NPs. Kuan et al. [102] offered an alternative method that included creating an electrochemical DNA biosensor for the detection of *Aphanomyces invadans* in fish by combining Au-NPs with a DNA reporter probe. With a lower detection threshold for fungal presence compared to PCR, this biosensor was designed for fish usage.

Fish viral disease diagnosis

A colorimetric test using Au-NPs and loop-mediated isothermal amplification (LAMP) was used to visually detect yellow head virus in shrimp by Jaroenram et al. [103]. Sensitivity, speed, and specificity were all much enhanced by this approach. Similarly, WSSV in shrimp was successfully detected utilizing a DNA-functionalized Au-NPs and LAMP combination, proving that the method is sensitive, accurate, and applicable in the field [104]. Using an Au-NPs-based biosensor for viral nucleic acid detection following RT-PCR amplification, Toubanaki et al. [105] devised a method for detecting nerve necrosis virus (NNV). Due to the lack of antibody conjugation, this method proved more cost-effective.





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Yang et al. [106] used magnetic NPs coated with rabbit anti-NNV antibody to conduct an immunomagnetic reduction experiment in grouper fish. The NNV was the target of the assay's development. Magnetic NPs' motion in response to an applied external magnetic field formed the basis of the immunodiagnosis. Virus antigen attachment resulted in cluster formation and reduced motility of antibody-coated NPs. The viral titer was calculated using a magnetic immunoassay analyzer. Unmodified Au-NPs were employed to develop a colorimetric test for detecting the spring viremia of carp virus (SVCV). In this method, Au-NPs were introduced after the SVCV probe. If the intended viral RNA was present, it hybridized with the probe, preventing it from holding the Au-NPs in place. The aggregation of Au-NPs resulted in a color shift from red to blue. In the absence of viral nucleic acid, the probe could freely bind to Au-NPs surfaces, preventing aggregation and maintaining the solution in a red color [107]. This approach demonstrated a high degree of specificity, rapidity, and did not require prior viral nucleic acid amplification. The same principle was applied to create a quick, accurate, and sensitive assay for detecting the DNA virus Cyprinid herpes virus-3 (CyHV-3) [108].

NANOTECHNOLGY-BASED FISH FEED

Nanomaterials have demonstrated superior effectiveness compared to bulk materials in enhancing the growth and health of cultured fish [109-112]. However, it is crucial to note that lower doses of nanomaterials can have detrimental effects. This review also explores the current applications of nanoparticles (NPs) in the nourishment, diagnosis, and treatment of fish pathogens [113]. Fish are conventionally fed through the administration of food pellets designed to meet their daily nutritional requirements, encompassing lipids, proteins, carbohydrates, minerals, and vitamins [114, 115]. Recent research has investigated the incorporation of nanomaterials as additives in fish meals to protect fish from antimicrobials. The implementation of modern techniques, such as rapid disease diagnostics, holds the potential to enhance the absorption of medications like hormones, vaccinations, and critical nutrients in cultivable organisms, revolutionizing the fisheries and aquaculture industries [115]. Aquaculture relies heavily on metal nanoparticles (NPs) like iron (Fe), iron oxide (FeO), selenium (Se), zinc (Zn), zinc oxide (ZnO), copper (Cu), and magnesium oxide (MgO). Studies have shown that young goldfish and sturgeon grow rapidly, by 30% and 24% respectively, when given iron nanoparticles. Crucian carp (Carassius auratus) also show better growth, antioxidant levels, and muscle selenium concentration when they eat different selenium sources, including nanoselenium and selenium methionine [115, 116]. Recent research suggests that post-larvae of freshwater prawns (Macrobrachium rosenbergii) have higher survival and growth rates when zinc, nano zinc, and copper are added to their diet [116-118]. Azolla microphylla-generated gold nanoparticles may considerably avert hepatic degeneration in Asian carp brought on by acetaminophen water pollution, according to research by Kunjiappan et al. [119]. Indicators of oxidative stress, hepatic ion levels, metabolic enzymes, hepatotoxic markers, abnormal liver histology, and altered tissue enzymes are all markedly improved by the presence of gold nanoparticles.

The Siberian sturgeon (Acipenser baerii) showed improvements in body composition, survival rate, and growth when exposed to Aloe vera based-nanoparticles, according to Sharif et al. [120]. Additionally, Cui et al. [121] noted that nanotechnology has been used to try to increase the retention time and bioavailability of naturally occurring bioactive chemicals. Encasing of curcumin in NPs has been shown using phospholipids [122], micelles [123,124] liposomes [125], hydrogels [126], and other methods [127]. In a recent study by Shah et al. [128], researchers were able to enhance the shelf-life of curcumin by encapsulating it in chitosan nanoparticles stabilized using Pickering emulsion. When dealing with hydrophobic bioactive substances, the Pickering emulsion method is regarded as the most effective way to encapsulate them and enhance their shelf-life. This is due to the solid particles (nutrients in this case) that stabilize the emulsion [129-131). There is less or no toxicity, higher repeatability and biocompatibility, easy and scalability manufacturing, and improved stability with these emulsions compared to traditional emulsions [132-134]. In addition, the composition and structure of these mixtures enhance the transportation of biological substances in the gastrointestinal system [135-138]. Presently, a variety of colloidal particles are employed in the manufacturing of Pickering emulsions, whereby solid particles aid in maintain the emulsion. Starch-based Pickering emulsion has been utilized to transport chemicals, such as hydrophobic antifungal agents [139-141]. There is promising research into the use of nano-crystalline, self-stabilized Pickering emulsions for drug administration and release, particularly with chemicals that are not very soluble [142]. In a recent study by Zhou et al. [143], cellulose nanocrystals were





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utilized to stabilize oil Pickering emulsions containing oregano essential oil. The purpose of this was to enhance the antibacterial effects of the emulsion against selected bacterial stains. In addition, a Zein/Arabic-gum NPs-maintained Pickering mixture was created using thymol to construct an antibiotic delivery system targeting E. col [144, 145]. In a recent study, Baldissera et al. [146,147] found that adding nerolidol-loaded nanospheres to the food of Nile tilapia infected with Streptococcus agalactiae resulted in decreased bacterial levels, improved survival rates, and protection against oxidative damage. Furthermore, nanoparticles may also be utilized to enhance the stability and bioavailability of dietary components, as well as to alter the physical characteristics of fish food. Even the use of modest amounts of nanoparticles can significantly improve the physical characteristics of food pellets.

As an illustration, when single-walled carbon nanotubes are added to the diets of trout, it leads to the formation of a solid pellet that remains intact while submerged in water. It is crucial to minimize contaminations and other food wastes in aqua forms caused by improper resilience, inadequate food stability, or unsuitable texture of the pellets. These factors result in substantial losses within the business [148]. Currently, there is significant research and investigation being conducted in the industry on the development of nano-formulations. An essential attribute of these systems is their versatility for many applications, including the administration of antibiotics, vaccines, medicines, and nutraceuticals, among others [149, 150]. Recent studies have concentrated on utilizing various biopolymers in the aquaculture industry [151-154]. The study discovered that when ascorbic acid is condensed in nanoparticles made from chitosan, it leads to an increase in the levels of ascorbic acid in the serum of rainbow trout when it is included in their diet. This increase in vitamin C levels also enhances the natural immune system, as indicated by higher amount of lysozyme and hemolytic serum complements, in comparison to the control groups receiving vitamin C and chitosan separately [155]. According to a recent study by Abd El-Naby et al. [156], feeding Nile tilapia (O. niloticus) with chitosan nanoparticle supplements can enhance their development and feed consumption. The authors contend that the presence of chitosan particles enhanced the activities of lipase and amylase. Furthermore, it has been shown that the utilization of these nanoparticles not only hinders the proliferation of all bacteria stains, but also enhances the inherent immune response.

In addition, another study have documented that the combination of ascorbic acid with chitosan NPs has an immunomodulatory impact on the toxicity induced organisms [157]. Furthermore, they highlight that the addition of this combinations leads to enhanced development and utilization of food in the presence of water contaminated with pesticides. And also, they highlight that the incorporation of chitosan NPs and ascorbic acid into the diet not only improves the antioxidant levels and generic immune response, but also potentially yields other beneficial health outcomes. These effects are evident in the cellular structure of hepatocytes and the overall well-being of *O. niloticus* when affected to sub-lethal levels of imidacloprid. In a recent finding, it was shown that the addition of dietary zeolites, either alone or in conjunction with chitosan NPs, can effectively alleviate the toxic impacts caused by introduction to imidacloprid [158]. However, it has also observed that the mixture of chitosan NPs with thymol can significantly enhance the development and usage of food in O. niloticus [159]. This is evident not only in the increased amount of lipases, catalases, and proteases, but also in the promotion of gastrointestinal villus length. However, the use of NPs as nutraceuticals in aquaculture (specifically shellfish and fish) for the purpose of enhancing value, reducing stress, and managing health is still in the early stages of consideration. The utilization of nutraceuticals is currently limited due to their high cost [160].

SUMMARY AND CONCLUSION

Nanomaterials have demonstrated superior effectiveness in improving the development and well-being of farmed fishes compared to larger materials. However, it is essential to ascertain the optimal particle size and dose to provide long-term advantages. Uncoated metal nanoparticles have been employed to contest microbial confrontation in aquaculture, but polymeric chitosan nanoparticles and PLGA-NPs have demonstrated efficacy in rapidly and cost-effectively identifying fish illnesses. Further investigation is required in the fields of nanocapsules, liposomes, dendrimers, and nanotubes to better understand fish illnesses. Additional research is crucial for the widespread use of nano technological methods in the development of vaccinations for fish. The aquaculture business has utilized





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nanotechnology to increase fish health management, include nanoscale components, and improve aquaculture feeds. Nevertheless, the majority of apps are now in their first phases, and the substantial expenses linked to them hinder sustainable expansion. Mineral nanoparticles outperform bulk materials as feed additives for improving fish growth and well-being. They are also more cost-effective and resource-efficient compared to bulk minerals. Higher levels of nanomaterials have been discovered to have detrimental impacts on the development and overall health of fish. Because of the encouraging results in these areas for the efficient diagnosis, treatment, growth, and development of fisheries, there is a pressing demand for further concentrated investigation into the use of nanoparticles in fish medicine and fish feed.

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Table, 1. Antimicrobial activity of nonoparticles against different microorganisms

Nanoparticals	Activity against the microorganisms	Reference
	Staphylococcus aureus, Edwardsiella tarda	[49]
	Pseudomonas fluorescens, Proteus species, and Flavobacterium species	[60]
	Vibrio harveyi	[36]
	Candida species; Streptomyces sp. VITPK1	[61,62, 63]
	HIV-1 virus	[64]
Ag-NPs (Chemically and	Bacterial strains: Streptococcus agalactiae, Aeromonas hydrophila and Vibrio alginolyticus	
Biologically synthesized)	fungal strains: Aspergillus flavus, Fusarium moniliforme and Candida albicans	[37]
Ag-Chitosan-NPs composite	H1N1 influenza A virus	[65, 66]
	Escherichia coli and Salmonella typhi	[69, 70]





Au-NPs	Candida species	[71, 72]
	Aermonas hydrophila, Edwardseilla tarda, Flavobacterium branchiophilum,	
	Citrobacter spp., Staphylococcus aureus, Vibrio species, Bacillus cereus, and	
	Pseudomonas aeruginosa	[49]
	Vibrio harveyi	[74]
ZnO-NPs	Pseudomonas aeruginosa, Escherichia coli, Enterococcus faecalis, Aspergillus	
ZHO-INI S	flavus, and Candida albicans	[39]
TiO ₂ -NPs	Streptococcus iniae, Edwardsiella tarda, and Photobacterium damselae	[75, 76]

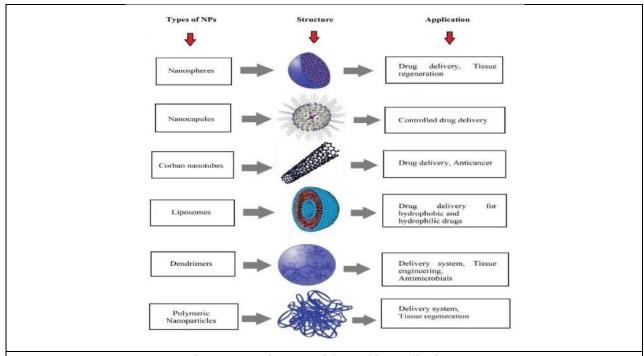


Fig. 1. Types of nanoparticles and its applications.

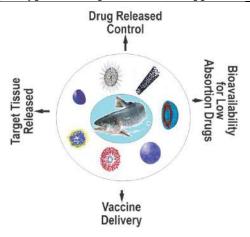


Fig. 2. Utilizing nanoparticals for drug delivery and targeted therapy in aquaculture offers numerous advantages





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RESEARCH ARTICLE

Exploring Competency Profiles among Emergency Department Nurses: **Factors Shaping Skills and Overcoming Barriers**

Suresh K1*, Thiruvenkadam Thiagarajan² and Ganesan Alias Kanagaraj Mariappan³

¹Assistant Professor, Department of Management Studies, SRM Valliammai Engineering College, (Affiliated to Anna University), Kancheepuram, Tamil Nadu, India.

²Associate Professor, Department of Management Studies, Sri Sivasubramaniya Nadar College of Engineering, (Affiliated to Anna University) Chennai, Tamil Nadu, India.

³Assistant Professor, Department of Management Studies, SRM Valliammai Engineering College, (Affiliated to Anna University) Chennai, Tamil Nadu, India.

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*Address for Correspondence Suresh K

Assistant Professor, Department of Management Studies, SRM Valliammai Engineering College, (Affiliated to Anna University), Kancheepuram, Tamil Nadu, India. Email: profsureshkkumar@gmail.com



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ABSTRACT

Gaining knowledge of and improving employee competencies is still essential for organizational success in the cutthroat healthcare industry of today. The study was conducted because comprehending employee competencies necessitates a targeted strategy to examine the essential competencies held by the personnel. The study aims to determine the level of competency of emergency department nurses, examine the abilities that the nurses possess in the largest and least amount, and identify the barriers that prevent the nurses from sharing their competency. An organized questionnaire was made and given to the respondents to collect the primary data required for the study. A simple random selection procedure was used to choose the responders. Utilizing factor analysis, reliability analysis, and correlation analysis, the data collected from 217 participants was tabulated in SPSS. To facilitate the application of the findings in improving emergency department nurses' competency for the benefit of both the individual and the organization, the research aims to bring out useful perspectives on the nurses' competency level to administrators, legislators, and human resource management practitioners.

Keywords: Competency Mapping, Human Resource Management, Skills, Health care, Nursing Competencies





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INTRODUCTION

The effectiveness of an organization is now evaluated based on how well it manages its most valuable resource—its people—and their capabilities rather than how well it uses material resources in the competitive world of today. Organizations now place a high focus on managing employee capabilities, and they have begun to make a concerted effort to identify the core competencies held by employees, their levels, and the steps that can be taken to enhance those competencies. An organization with highly skilled personnel enjoys higher advantages in the current business environment. Competence is an underlying quality of a person that includes their motivations, characteristics, abilities, components of their image or social role, and information that they may apply on a personal level. It is the ability to act with authority and highly qualified awareness; it is the capacity to perform at the highest level in terms of training, expertise, and knowledge. The combination of an individual's aptitudes, personality, and experience tailored to a role or task in the current and emerging context, which accounts for sustained success within the confines of company principles. Competency is the set of success characteristics required to accomplish significant goals in a certain job or work capacity inside a specific organization. Healthcare systems nowadays deal with several difficulties. Increasing healthcare demands and expenses are a consequence of an aging population with numerous pathologies and non communicable diseases, which are made worse by an explosion in new medical technologies.

The maldistribution and scarcity of labor, as well as the inadequate quality and safety of services provided, exacerbate these issues further. The provision of health education by nurses in acute and community settings is widely acknowledged as a critical role that they play in improving favorable health results and the general effectiveness of the provision of healthcare. Since they are the most approachable medical staff in this setting and have frequent interaction with patients, hospital nurses in particular play a significant role in routine health education practices. Crucially, hospitalization affords a "window of opportunity" to promote lifestyle changes because of the numerous teaching opportunities this setting presents. Hospital nurses have been seen to express challenges in carrying out everyday initiatives in health education, and it is possible that they lack confidence in their ability to deliver effective health education. Interventions designed to improve nurses' health education competency should take into account contextual elements such having organizational support and be customized to the nurses' individual learning requirements and characteristics. The required level of performance for nurses includes judgment, knowledge, skills, and talents. Therefore, it is appropriate to regard as competent any individual who performs well at the expected level of nursing education and orientation. (Alshammari et al., 2022)Competencies are separated into two categories by the Emergency Nurses Association Nurse Practitioner Team: special and general.

A few examples of common clinical competencies include the ability to operate in a team, following professional standards, having a basic understanding of medicine, and having basic cultural traits and professional principles. To accomplish the duties of the job, a nurse needs to have certain basic abilities. Their clinical knowledge is essential, particularly in the emergency department where they treat injured patients all day long and make up the majority of the medical staff. Emergency Department (ED) nurses, for instance, are expected to acquire the capacity to predict patients' future situations and take appropriate action. In a hectic environment, emergency department nurses must recognize, rank, and assess cases that pose a threat to life, administer both emergency and non-emergent care in accordance with equitable standards, and demonstrate expertise and high-quality care. Continuously evaluating the clinical competency of ED nurses is necessary to identify areas that require improvement and promotion in order to provide consistently high-quality treatment. This aids ED nurses in confirming the suitability of the care they provide, as well as the effectiveness and caliber of it. For this reason, an evaluation of ED nurses' competency is necessary. Jeon et al. state that competency evaluation is utilized to advance current methods and encourage lifelong learning. Understanding the nursing competency development process is therefore essential for ongoing professional development. It is considered that nurses are very autonomous and independent enough to act on their own initiative without oversight. As a result, they are skillfully providing the patient, their family, and other significant individuals with the knowledge and emotional support required to handle the circumstance. Therefore, to provide





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safe, efficient, and professional nursing care, a competent individual needs to possess these attributes, be driven to use them, and be able to do so effectively. To lay the foundation for future developments in nursing competency in the emergency department, a precise definition of nurse competency is required. Unfortunately, there is still a lack of consensus on what constitutes a competent nurse, even though nursing care can only be improved with skilled nurses. This implies that issues with defining and proving nursing competency, determining what proficiency levels are required of nursing practitioners, and creating educational resources will never go away. Moreover, the environment has a significant influence on the rate at which nurses become competent. Determining talents is therefore essential to enhancing nursing care. Therefore, mapping the ED nurse competencies and the related factors that impact those competencies is the aim of this research. additionally, to identify obstacles that limit their potential.

REVIEW OF LITERATURE

Studies have shown that nurses' competency has received a great deal more attention throughout time and is now a worry for healthcare professionals. Numerous scholars have previously investigated the elements that affect nursing practitioners' ability. Since it would be very helpful in defining the research objectives, a thorough review of the prior literature in the field of employee competency has been done. The study sought to characterize the differences in perceptions among clinical nurses on the impact of an alternative supervision model on their ability to evaluate nursing students during clinical practice. The research employs a phenomenographic approach in a qualitative and descriptive design. 49 clinical nurses from five different nursing homes served as informants. Three categories—"pressure," "encouragement," and "development"—are used to characterize the experiences of the clinical nurses. The clinical nurses' evaluation of the nursing students and their function as educators were aided by the alternative supervision approach(Struksnes et al., 2012). The systematic review (Belita et al., 2018) gives an up-to-date understanding of the evidence on evidence-informed decision-making (EIDM) competence assessments in nursing. This will help identify measures that may be useful and reliable for use in various nursing practice settings.

Developed in collaboration with a Health Sciences Librarian, the search strategy makes use of internet databases, expert advice, manual reference list searches, key journals, websites, conference proceedings, and grey literature. The study looked at the literature on competency structure, components, assessment, and nursing competency definitions and characteristics in Japan. The study also examined methods for nurse competency training. The idea of nursing competency has not yet reached its full potential, despite the fact that skills are crucial for raising the standard of care provided by nurses. As a result, there are still problems with training curriculum development, defining and organizing nursing competency, and figuring out what competencies nursing practitioners need to have(Fukada, 2018). The study conducted by (Nilsson, Engström, Florin, Gardulf, & Carlsson, 2018) with an objective to create a condensed version of the NPC Scale and assess its construct validity and internal consistency. A sample of 1810 nursing students who were about to graduate from 12 Swedish universities served as the basis for the study. The 35-item NPC Scale Short Form (NPC Scale-SF) produced encouraging results, with a six-factor structure accounting for 53.6% of the variance in the total. When combined with other tools, this 35-item scale can be a useful tool for doing more in-depth evaluations of nursing students' and registered nurses' self-reported competence.

The goal to assess clinical competency among critical care nurses in Kermanshah, Iran, (Faraji, Karimi, Azizi, Janatolmakan, & Khatony, 2019) studied the relevant demographic characteristics. Using stratified random sample, 155 Iranian nurses were chosen for this cross-sectional investigation. The "Nurse Competence Scale" and a questionnaire for personal information were among the data collection instruments. The results of the study showed that clinical competence was used at a "good level" in real practice and that critical care nurses in Kermanshah had a "very good" rating for it. Given the importance of clinical competencies in nursing practice, it is critical to carry out an objective evaluation of nurses' clinical competence and to put proactive measures in place that promote the application of their knowledge. The study's by (Immonen et al., 2019) was focused towards the goal to identify the most up-to-date data that might be used to assess nursing students' clinical competency. The electronic databases CINAHL, PubMed, Eric, Medic, and the JBI Database of Systematic Reviews and Implementation Reports were





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searched in the fall of 2018. Six reviews made it in after being critically evaluated. Assessment tools used to gauge students' nursing competency often cover the following topics: reason, communication and interpersonal relationships, nursing procedures, ethical practices, professional attributes, and critical thinking. Students' education is guided and vital support networks are provided by mentoring and clinical learning environments. The development of standardized and methodical assessment procedures as well as the utilization of trustworthy and legitimate equipment are still necessary.(Jao, Chen, & Sun, 2020)studied the path links influencing students' learning outcomes in the clinical practicum using LISREL software. Participants were 392 senior students from two nursing programs in central Taiwan who had completed their last internship. The Competency Inventory of Nursing Students, the Teaching Competence of Nurse Preceptor questionnaire, the Student Evaluation of the Clinical Education Environment, the Level of Reflective Thinking, and the Metacognitive Inventory for Nursing Students are some of the structured questionnaires that need to be distributed and filled out. The results suggest that metacognition and reflection, by teaching competency, may influence nursing competence indirectly.

The clinical learning environment can have a direct impact on nursing competency, while metacognition can have an indirect impact. Reflection and metacognition are essential for the development of nursing competence in clinical settings(Jao et al., 2020). During the COVID-19 pandemic in the spring of 2020(Konrad, Fitzgerald, & Deckers, 2021) presented a difficulty to numerous nursing schools, forcing them to switch from in-person to online instruction. In this article, educational concepts for an online course on the principles of nursing that enhance clinical competency are discussed. Presentations are made regarding faculty planning, teaching pedagogy used throughout this shift, and regulatory considerations. The writers also go into potential ramifications and ideas for achieving learning goals for an online healthcare course. The study conducted by (Alshammari et al., 2022)was focused towards mapping ED nurses' competencies, barriers, and influencing factors is the goal of the study. This study used cross-sectional methodology and was carried out at Hail's government hospitals, including the hospitals in the villages. The 227 ED nurses who were selected by convenience sampling were the study's participants. To get the data, the researchers used a Google Form survey. The study came to the conclusion that ED nurses possessed a high level of competence, particularly in clinical care leadership. Based on the study's findings, it is recommended that ED nurses' competency be continuously measured. The ED requires training needs assessments, which should be carried out on a regular basis and provide ongoing nursing education tailored to nurses' needs. The aim of the research (Suresh, 2020) is to ascertain the degree of competences acquired by employees and to evaluate which competencies are most and least prevalent. The items in the questionnaire used to gauge the respondents' competencies were framed with assistance from past research in the field of competency mapping.

Subjects and Methods

A cross-sectional methodology was used in this study to map out the competencies of ED nurses as well as pertinent variables and obstacles to learning them. The 217 ED nurses employed in government and private hospitals, including those in villages, were the study's participants. The study comprised ED nurses who met the following criteria: (a) were willing to participate; (b) could understand English; and (c) had worked in the ED for at least six months. The exception applied to nurses who moved to the emergency department to substitute for absentee nurses. Convenience sampling was employed by the researchers, bearing in mind the inclusion criteria. With 217 participants (73% response rate), the sample size calculator was used online at https://www.calculator.net/sample-size-calculator. The true value had a 95% confidence level and was within ± 5% of the measured/surveyed value.

Objectives and Limitations

The broad objective of the research is to study and assess the competency of Emergency Department Nurses employed in government and private hospitals in Chennai. The purpose of this study is to map ED nurses' competences and the related variables that affect them. This study also attempts to identify the obstacles that impact the competences of ED nurses. We must consider the limitations of this study. One of the fundamental limitations of cross-sectional studies, for example, is that it is impossible to identify the intricate relationship between the exposure and the outcome because they are viewed concurrently. Furthermore, leaving out nurses—for example, those who are illiterate in English—might not accurately reflect the entirety of the situation when evaluating ED competencies.



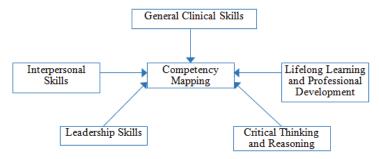


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As such, care must be taken when analyzing the data. More participants and a bigger study population are recommended in order to accurately map the competences and generalize the contributing components affecting the competencies.

Conceptual Framework

The conceptual framework of the study has been established by the researcher in accordance with the literature review and taking the research objectives into consideration. The Competency Inventory for Registered Nurses (CIRN)(Alshammari et al., 2022)(Jao et al., 2020), which served as the foundation for the study's questionnaire, was created for this research. The following five categories comprised the 25 questions that made up the questionnaire: Leadership, Critical Thinking and Reasoning, Lifelong Learning, Interpersonal Skills, General Clinical Skills, and Professional Development



RESEARCH DESIGN

Descriptive research design was used for the study. In Chennai, Tamil Nadu, India, nurses working in emergency rooms of both public and private hospitals provided the primary data. Because it makes it easier to gather data from a wide population, the survey method was suitable for the study. To conduct the investigation, a straightforward random sample strategy was used. To gauge the degree of skills, a systematic questionnaire was created in accordance with the study's goals. Each skill, which had five statements each, was scored on a 5-point Likert scale, ranging from 5-Strongly Agree to 1-Strongly Disagree.A straightforward random sampling technique was used to gather the samples. There were 300 survey questionnaires distributed in all, of which 217 complete and usable surveys were received. SPSS 20.0 was used to tabulate and analyze the data that was gathered from the respondents. Among the statistical tests used to assess and draw conclusions from the study were Anova, Correlation, T Test, Factor Analysis, and Reliability Analysis.

RESULTS AND DISCUSSION

The goal of this study is to map the competences of emergency department nurses along with the related factors that impact those competencies. Additionally, this study attempts to identify the obstacles that impact the competences of ED nurses. All things considered; emergency department nurses gave themselves a very high competence rating. This indicates that nurses are aware of the necessity to prove their clinical competence to carry out tasks and support the organization's objectives. It has been determined that the greatest competencies are in clinical care, leadership, and interpersonal skills. The findings of this study corroborate those of an earlier study by V and Tamadoni et al., in which nurses assessed their level of competence in clinical care provision, leadership, and legal/ethical practice as being extremely high. In fact, it is considered that nurses must be competent professionals with leadership skills despite their intimate contact with patients and healthcare institutions. Leaders can identify inefficiencies in procedures, policies, practices, organizational structures, and processes that impede the provision of optimal patient care. Furthermore, it is acknowledged that a nurse's leadership qualities have a big influence on patient outcomes and the hospital atmosphere. The table above (Table 1) includes the demographic profile of the respondents. It can be





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deduced from the preceding table that 145 (66.8%) of the respondents are female and 72 (33.2%) are male. Seventyseven (35.3%) of the respondents are in the 20-29 age group; sixty-two (28.6%) are in the 30-39 age group; fortyforty-nine is represented by 63 (29.0%) of the respondents; and fifty-fifty-nine is represented by 15 (6.9%) of the respondents. In terms of experience, 122 of the respondents (56.2%) are of 5 years and above years of experience in the Emergency Department; in the salary category 106(48.8%) have reported their earning to be between 25001 – 50000 per month. Table 2 shows that the item "I think I'm capable of being a leader since I like to set objectives and work toward achieving them." has the lowest mean value. It hovers close to neutral. The item "Using the information at my disposal, I determine if a patient needs nursing intervention" (category Agree) had the highest mean rating. Based on a thorough study of the data provided, it can be deduced from the table that most respondents are confident in their ability to offer nursing care to patients, and they all concur that they possess general clinical skills. To identify the pertinent variables that define each employee's skill level, a factor analysis was performed. The results of the Bartlett's test of sphericity values and the Kaiser-Meyer-Oklin (KMO) tests are displayed in Table 3. The KMO measure of sampling adequacy value was determined to be 852 for the full sample. Bartlett's test of sphericity (p=0.000) shows that the correlation between the variables is statistically significant. The KMO and Bartlett's test findings indicated that the data was appropriate for factor analysis. Principal Component Analysis and the Varimax rotation approach were used in the investigation. The communalities of the loaded items and the percentage of variation explained by each item in the study, which varies from 50.5 to 75.4, are shown in Table 4. The six components that were recovered and accounted for 64.68 percent of the variance in the total sample are shown by the Principal Component Analysis (Table 5). The PCA's scree plot is displayed below. To evaluate the dependability of the factors extracted, a reliability test was carried out. The reliability was assessed by calculating the internal consistency Cronbach's Alpha coefficient (Table 7). The criteria were shown to be extremely dependable in predicting the Competencies, as indicated by the attained Alpha coefficient of 910.

Table 8 displays the values of the reliability coefficients for the factors. (Wang & Shi, 2011)According to Nunnally (1978), an alpha of at least 0.600 was sufficient for preliminary study. Cronbach's alpha estimate of 0.723 has been estimated for Leadership, 0.828 for General Clinical Skills, 0.700 for Interpersonal Skills, 0.763 for Critical Thinking and Reasoning, and 0.723 for Lifelong Learning and Professional Development. Given that every study's Cronbach's alpha was significantly. One statistical evaluation method that can be used to assess the strength of a link between two numerically measured continuous variables is correlation analysis. was conducted to investigate the relationship. The table below displays the correlation study's findings. Based on the table below, one may deduce that there is a perfect correlation and a substantial relationship between the variables. The study also intends to identify the barriers that affect ED nurses' competencies. Research indicates that those with ED training possess superior abilities compared to those without any training prior to entering the ED. It follows that nurses with ED training get their skills from their education. Because licensed providers are legally required to guarantee that each employee has completed a comprehensive induction and obtained a basic, appropriate level of training, staff training is essential. It is advised that in order to create training methods that cover the competency development sequence, it is crucial to understand the traits and elements of these nursing skills as facets of the clinical leader. To assess nurse competency and study the outcomes of competency improvement, more research is needed. The training does not correspond with the needs of emergency department nurses, which is one of the obstacles to gaining competence. The reason for this is because no evaluation of the training needs was conducted prior to the training or course being delivered.

CONCLUSION

An organization's core competences are identified as part of the mapping process, and these competencies are then attempted to be integrated throughout all of the organization's activities. Mapping the organizational abilities needed and putting policies in place to help employees recognize and acquire these competencies are important tasks performed by the human resources department. The research outcomes reveal that the emergency department





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nurses demonstrated exceptional competency, particularly in the area of clinical care leadership. Their proficiency was influenced by their older age group, less than five years of ED experience, and training. The absence of support from leaders and the training's incompatibility with ED nurses' needs were obstacles. The results of the study indicate that ongoing competency assessment is necessary to raise ED nurses' proficiency. Particularly in the ED, training needs assessments should be carried out on a regular basis to provide ongoing nursing education based on the needs of the nurses. The study has offered a road map for comprehending the skills emergency department nurses in Chennai's public and private hospitals possess. The study's findings can have a greater managerial influence and offer HR managers, directors of departments and units, and policy makers a blueprint for developing competency development initiatives that will support the promotion of the essential skills. In order to create a solid competency-based HR framework, the study offers a foundation for organizing and expediting all HR tasks, such as talent mapping, hiring, on boarding, development, and performance appraisal.

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Table 1: Table showing the demography of the respondents

Factor		%	Count	Valid %	Cumulative %
Condor	Female	145	66.8	66.8	66.8
Gender	Male	72	33.2	33.2	100
	20-29	77	35.5	35.5	35.5
A	30-39	62	28.6	28.6	64.1
Age	40-49	63	29	29	93.1
	50-59	15	6.9	6.9	100
Marital Status	married	129	59.4	59.4	59.4
Maritai Status	Unmarried	88	40.6	40.6	100
	5 Yrs. and above	122	56.2	56.2	56.2
E	3 – 4 Years	56	25.8	25.8	82
Experience in Emergency Department	2 – 3 Years	31	14.3	14.3	96.3
	1 – 2 Years	8	3.7	3.7	100
	Up to25000	69	31.8	31.8	31.8
Salary	25001-50000	106	48.8	48.8	80.6
_	50001-75000	42	19.4	19.4	100

Table 2: Table showing the mean and the standard deviation

Table 2: Table snowing the mean and the standard deviation	Mean	Std. Deviation	Analysis N
Using the information at my disposal, I determine if a patient needs nursing intervention.	3.94	1.323	217
I administer nursing care to patients based on their individual needs.	3.95	1.301	217
I try to learn as much as I can about the patient so that I can create the most effective nursing plan.	3.76	1.268	217
I prioritize nursing duties according to the demands of my patients.	3.84	1.290	217
I make an effort to give patients thorough follow-up treatment that is tailored to their needs.	3.65	1.409	217
I think I'm capable of being a leader since I like to set objectives and work toward achieving them.	3.49	1.371	217
I see myself as being helpful to all of my peers, and I believe that I accept full responsibility for the work that I am doing.	3.80	1.383	217
Guidance without coercion is essential to effective leadership.	3.79	1.444	217
I take pleasure in working in teams.	3.61	1.433	217
I administer nursing care to patients based on their individual needs.	3.78	1.360	217
I try to learn as much as I can about the patient so that I can create the most effective nursing plan.	3.74	1.330	217
I often look to others for validation and encouragement.	3.61	1.357	217
I make a special effort to uplift those in the group, and I pay close attention to what they have to say.	3.72	1.515	217
To ensure that people don't squander time or circle around in circles, I push for action.	3.77	1.316	217
Every time a patient's condition changes, I attempt to identify the underlying cause.	3.64	1.395	217
I make an effort to examine patients' issues from several perspectives.	3.51	1.320	217
I carefully evaluate the patient's situation before drawing logical conclusions.	3.85	1.406	217





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I consider whether there is sufficient empirical evidence to warrant nursing interventions every time.	3.46	1.430	217
I frequently turn to other people for approval and inspiration.	3.76	1.305	217
I prefer to weigh all of my options before making a decision, and I am aware of where to get educational materials and how to use them.	3.70	1.374	217
I know exactly what I still need to learn.	3.51	1.388	217
I like to look for answers to things.	3.59	1.479	217
I use other resources and technologies to further my education.	3.35	1.521	217
I manage my time well at work and establish learning objectives.	3.81	1.589	217
I frequently turn to other people for approval and inspiration.	3.78	1.290	217
Valid N (listwise)	217		

Table 3: KMO and Bartlett's Test

Kaiser-Meyer-Olkin Measure	.852	
	Approx. Chi-Square	2193.052
Bartlett's Test of Sphericity	df	300
	Sig.	.000

Table 4: Communalities

	Initial	Extraction
Using the information at my disposal, I determine if a patient needs nursing intervention.	1.000	.709
I administer nursing care to patients based on their individual needs.	1.000	.632
I try to learn as much as I can about the patient so that I can create the most effective nursing plan.	1.000	.610
I prioritize nursing duties according to the demands of my patients.	1.000	.667
I make an effort to give patients thorough follow-up treatment that is tailored to their needs.	1.000	.618
I think I'm capable of being a leader since I like to set objectives and work toward achieving them.	1.000	.620
I see myself as being helpful to all of my peers, and I believe that I accept full responsibility for the work that I am doing.	1.000	.740
Guidance without coercion is essential to effective leadership.	1.000	.695
I take pleasure in working in teams.	1.000	.754
I administer nursing care to patients based on their individual needs.	1.000	.727
I try to learn as much as I can about the patient so that I can create the most effective nursing plan.	1.000	.654
I often look to others for validation and encouragement.	1.000	.612
I make a special effort to uplift those in the group, and I pay close attention to what they have to say.	1.000	.665
To ensure that people don't squander time or circle around in circles, I push for action.	1.000	.576
Every time a patient's condition changes, I attempt to identify the underlying cause.	1.000	.676
I make an effort to examine patients' issues from several perspectives.	1.000	.630
I carefully evaluate the patient's situation before drawing logical conclusions.	1.000	.631
I consider whether there is sufficient empirical evidence to warrant nursing interventions every time.	1.000	.644
I frequently turn to other people for approval and inspiration.	1.000	.505
I prefer to weigh all of my options before making a decision, and I am aware of where to get educational materials and how to use them.	1.000	.585
I know exactly what I still need to learn.	1.000	.606





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I like to look for answers to things.	1.000	.618
I use other resources and technologies to further my education.	1.000	.712
I manage my time well at work and establish learning objectives.	1.000	.575
I frequently turn to other people for approval and inspiration.	1.000	.708
Extraction Method: Principal Component Analysis.		

Table 5: Component Matrix^a

	Component						
	1	2	3	4	5	6	7
Using the information at my disposal, I determine	(1)	-					
if a patient needs nursing intervention.	.616	.483					
I administer nursing care to patients based on their	.561	-					
individual needs.	.501	.428					
I try to learn as much as I can about the patient so that I	.636	-					
can create the most effective nursing plan.	.000	.415					
I prioritize nursing duties according to the demands of	.592	-					
my patients.		.381					
I make an effort to give patients thorough follow-up	.568	-					
treatment that is tailored to their needs.		.436					
I think I'm capable of being a leader since I like to set	.502	.498					
objectives and work toward achieving them. I see myself as being helpful to all of my peers, and I		.490					
believe that I accept full responsibility for the work that I	.565		-		.344		
am doing.	.505		.365		.544		
Guidance without coercion is essential to effective			_				
leadership.	.591	.306	.371				
I take pleasure in working in teams.	.543				.501		.339
I administer nursing care to patients based on their	(1)		-				
individual needs.	.616		.444				
I try to learn as much as I can about the patient so that I	.541	.384			.382		
can create the most effective nursing plan.	.541	.504			.502		
I often look to others for validation and encouragement.	.566		-				
			.515				
I make a special effort to uplift those in the group, and I	.436	.303	.360	.422			
pay close attention to what they have to say.							
To ensure that people don't squander time or circle around in circles, I push for action.	.578						
Every time a patient's condition changes, I attempt to							
identify the underlying cause.	.482			.424		.379	
I make an effort to examine patients' issues from several							
perspectives.	.609						.420
I carefully evaluate the patient's situation before				220			
drawing logical conclusions.	.576			.330			
I consider whether there is sufficient empirical evidence	600				-		
to warrant nursing interventions every time.	.609				.382		
I frequently turn to other people for approval and	.588			-			
inspiration.	.500	<u> </u>		.326			





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I prefer to weigh all of my options before making a decision, and I am aware of where to get educational materials and how to use them.	.627				
I know exactly what I still need to learn.	.521		- .364		- .341
I like to look for answers to things.	.599		- .352		
I use other resources and technologies to further my education.	.451		.452	.432	
I manage my time well at work and establish learning objectives.	.515	.428			
I frequently turn to other people for approval and inspiration.	.472	.312		.481	.337
Extraction Method: Principal Component Analysis.			·		
a. 7 components extracted.					

Table 6: Total Variance Explained - Extraction Method: Principal Component Analysis.

Compone		Initial Eigenvalues Extraction Sums of Squared Rotation Sums of Squared					-		
nt					Loadings			Loading	gs
	Tot al	% of Varian ce	Cumulati ve %	Tot al	% of Varian ce	Cumulati ve %	Tot al	% of Varian ce	Cumulati ve %
1	7.87 1	31.482	31.482	7.87 1	31.482	31.482	3.41 2	13.647	13.647
2	1.91 2	7.647	39.129	1.91 2	7.647	39.129	2.86 9	11.474	25.121
3	1.58 7	6.348	45.477	1.58 7	6.348	45.477	2.32 4	9.294	34.415
4	1.47 5	5.900	51.378	1.47 5	5.900	51.378	2.07 5	8.301	42.716
5	1.24 2	4.969	56.347	1.24 2	4.969	56.347	2.01	8.054	50.770
6	1.07 0	4.281	60.628	1.07 0	4.281	60.628	1.86 4	7.456	58.225
7	1.01 4	4.055	64.683	1.014	4.055	64.683	1.614	6.457	64.683
8	.869	3.477	68.160						
9	.832	3.328	71.488						
10	.744	2.977	74.466						
11	.729	2.915	77.381						
12	.669	2.677	80.058						
13	.586	2.344	82.402						
14	.537	2.150	84.552						
15	.502	2.006	86.558						
16	.467	1.866	88.424						
17	.453	1.813	90.237						
18	.438	1.751	91.988						
19	.373	1.492	93.480						





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20	.344	1.375	94.854			
21	.314	1.256	96.110			
22	.288	1.150	97.260			
23	.248	.991	98.251			
24	.234	.936	99.187			
25	.203	.813	100.000			

Table 8: Table showing the Reliability Statistics

Cronbach's Alpha	N of Items
.892	25

Table 8.a: Table showing the Cronbach's Alpha Value

Factor	Cronbach's Alpha	N of Items
Leadership	.723	5
General Clinical Skills	.828	5
Interpersonal Skills	.700	5
Critical Thinking and Reasoning	.763	5
Lifelong Learning and Professional Development	.723	5

Table 9: Correlations

Table 9: Correlati		General Clinical Skills	Leadership	Interpersonal Skills	Critical Thinking and Reasoning	Lifelong Learning and Professional Development
General Clinical Skills	Pearson Correlation	1	.544**	.473**	.539**	.484**
	Sig. (2- tailed)		.000	.000	.000	.000
Leadership	Pearson Correlation	.544**	1	.613**	.602**	.493**
	Sig. (2- tailed)	.000		.000	.000	.000
Interpersonal Skills	Pearson Correlation	.473**	.613**	1	.602**	.459**
	Sig. (2- tailed)	.000	.000		.000	.000
Critical Thinking and Reasoning	Pearson Correlation	.539**	.602**	.602**	1	.591**
	Sig. (2- tailed)	.000	.000	.000		.000
Lifelong Learning and Professional Development	Pearson Correlation	.484**	.493**	.459**	.591**	1
	Sig. (2- tailed)	.000	.000	.000	.000	
Learning and Professional	Pearson Correlation Sig. (2- tailed)	.484**	.000	.459**	.000	





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RESEARCH ARTICLE

Revitalize Through Technology: How Online Shopping Habits of Youth **Shape Family Dynamics**

Vani Haridasan^{1*}, Kavitha muthukumaran^{1*} Vidyaa Lakshmi P² and Srinithi JP²

¹Associate Professor, Department of Management Studies, Sri Sivasubramaniya Nadar College of Engineering, Chennai, Tamil Nadu, India.

²Student, Department of Management Studies, Sri Sivasubramaniya Nadar College of Engineering, Chennai, Tamil Nadu, India.

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*Address for Correspondence

Vani Haridasan, Kavitha muthukumaran Associate Professor, Department of Management Studies, Sri Sivasubramaniya Nadar College of Engineering, Chennai, Tamil Nadu, India.



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ABSTRACT

This research investigates the profound impact of youth's online buying behavior on family purchasing decisions, spanning across generations from Gen Z to Gen X and Baby Boomers. Our objective is to explore the extent to which the younger generation influences the buying decisions of older family members, particularly towards online channels. Employing a mixed-methods approach, conducted a cross-sectional surveys and interviews to comprehensively capture insights into this phenomenon. By targeting Gen Z and older individuals, we aim to analyze the influence of tech-savvy youth on traditional consumer segments. We hypothesize a connection between youth's online purchases and older family members' buying patterns. The era of pandemic has triggered a higher-level comfort and elevation in the e commerce field. The new age on retail is evidently welcomed by the youth of today and the easy shop access is widely accepted by various age groups who once where traditional on-store vendee meanwhile the offline only retailers are affected drastically.

Keywords: Online Purchase, Youth Influence, Family Buying decision, Affecting Retailers

INTRODUCTION

The evolution of E- market has grown tremendously and the graphical movement from physical to electronic mode is so vivid. This research focuses on the semi urban take in the dramatic shift of the buyer's environment. This research delves into the intricate relationships between generations and their influence on purchasing decisions,





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particularly in the context of online buying behavior, it is crucial to explore the interplay between the tech-savvy youth and older family members, spanning generations from the digitally native Gen Z to the more traditional Gen X and Baby Boomers. Our primary goal is to assess the degree to which youth influence family members' buying decisions towards online channels. To achieve this, we employ a mixed-methods approach, utilizing cross-sectional surveys and interviews. This approach allows us to not only quantify trends but also gain qualitative insights into the motivations and attitudes driving these behaviors. The research hypothesis posits a connection between the online purchasing patterns of the youth and the buying behaviors exhibited by older family members. By targeting specific age groups, we aim to discern the nuances in the influence wielded by the younger generation on traditional consumer segments. Furthermore, we explore the transformative journey of the e-commerce market, emphasizing the pronounced impact of the pandemic on consumer behavior. The era of lockdowns and social distancing has propelled a higher level of comfort and elevation in the e-commerce field. The COVID-19 pandemic has accelerated this shift, triggering heightened comfort and reliance on e-commerce platforms. This research underscores the impact of these changes on both online and offline retailers, particularly emphasizing the challenges faced by traditional, offline-only retailers. As we embark on this exploration, we seek to contribute valuable insights for businesses, marketers, and policymakers navigating the ever-changing terrain of consumer preferences and retail dynamics.

OBJECTIVES

- To examine the buying behaviour of youth.
- To study the significance of online purchasing power of youth and its influence.
- To analyse the shift in the mindsets of family members and how it affects the retail shops.

NEED OF THE STUDY

This study is essential for retail shoppers to grasp the evolving consumer landscape. It offers insights into the increasing shift towards online shopping, enabling retailers to adapt their strategies effectively. Understanding intergenerational influences allows retailers to tailor marketing approaches for both tech-savvy youth and older consumers. With the pandemic accelerating the move to e-commerce, enhancing digital presence becomes a priority. Exploring semi-urban areas provides a nuanced understanding for customization, while acknowledging challenges faced by offline-only retailers guides the development of survival strategies, potentially involving digital integration or a hybrid model to stay competitive.

PROBLEM STATEMENT

- The tastes and preferences of the consumers are changing at a rapid rate. The marketers are finding it difficult to cope up with the changing need of the customers.
- E-com operators targeting youth and using them as a key indicator for their marketing strategy and this is directly affecting the Traditional Retail operators.

LITERATURE REVIEW

Ramya N, Dr S.A.Mohamed Ali (2016) delve into Consumer Buying behaviour by identifying and understanding the factors that influence their customers, brands can develop a strategy, a marketing message (Unique Value Proposition) and advertising campaigns more efficient and more in line with the needs and ways of thinking of their target consumers, a real asset to better meet the needs of its customers and increase sales. Lastly, the consumer analyses the prevailing prices of commodities and takes the decision about the commodities he should consume. Praveen K.S. Naik and Shivalinge Gowda M (2017). This study examines the factors influencing the online buying behaviour of the youth particularly the teenage groups. Convenience sampling method was used to select the sample of 30 teenage college students and a self-administered questionnaire was used to obtain the data. The study revealed some important factors influencing online shopping such as availability, low price, promotions, comparison, convenience, customer service, perceived ease of use, attitude, time consciousness, trust and variety seeking. Ritwik





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Maity, Sukjeet Kaur Sandhu (2021), Stated that their research is practically explored in embracing an online buying network from the customer. This research addresses many significant research issues pertaining to the acceptability of the user to choose online shopping portals while buying items. Such topics were divided into four variables i.e. societal acceptance of online purchasing, product preference for online purchasing, product variety availability for online purchasing and lastly convenience of online purchasing. As a result, a framework is needed to coordinate a complex process for the effects of these factors. Indrila Goswami Varma, Ms. Rupa Agarwal (2014) This paper aims to identify the factors that affect the online buying behaviour of women particularly homemakers in Western suburbs of Mumbai .Educated , urban homemakers form a significant prospect for traditional retailers. The question that arises is whether this segment is also buying online or has remained untapped by the e-retailers? Research has examined the role of different factors on individuals' ecommerce adoption, such as geography and store accessibility, perceived risk and online shopping benefits, typology of online stores, enjoyment and trust in Web sites , gender differences in attitudes toward online shopping ,and impact of consumers socio-economic conditions. Halima Shebuge (2019) The focus of this study was to analyse factors that influence youths' online buying behaviour in Tanzania. Specifically, the study aimed at finding out satisfaction levels, factors motivating youths to purchase online and the challenges that are facing them. The findings of the study revealed that large number of respondents were satisfied with sales services that are being provided in local and international online platforms in Tanzania, due to the comparison capacity, goods customization, and ability to trace orders. The study thus recommends establishing many local online market places to reach many customers together with laws governing online businesses.

METHODOLOGY

This is a Mixed-methods (i.e. both Quantitative and Qualitative analysis) research design to investigate online purchasing behaviors among two distinct age cohorts: Generation Z and mid-age individuals. The study included a sample of 224 responses from Generation Z and 135 responses from the mid-age group.

CONCEPTUAL FRAMEWORK TO UNDERSTAND THE FACTORS THAT INFLUCENCE FAMILY MEMBERS TO CHOOSE ONLINE PLATFORM DUE TO GEN Z PURCHASING BEHAVIOUR DATA COLLECTION

Surveys were administered to gather responses, focusing on various aspects of online purchasing decisions, including recommendations, assistance-seeking, and the impact on family spending habits. The survey instrument was designed to provide nuanced insights into factors influencing buying patterns and preferences.

ANALYSIS AND FINDINGS

Responses from both age groups underwent detailed analysis using Microsoft Excel. Statistical methods included correlation analysis, regression modeling, chi-square testing, and ANOVA. These analyses were instrumental in uncovering relationships, patterns, and trends within each age cohort.

Correlation between Recommending Products and Purchasing Decisions:

- The analysis revealed a strong positive correlation between the likelihood of recommending products or online stores to family members and their subsequent purchasing decisions.
- This suggests that when younger family members, especially those belonging to Gen Z, recommend products
 or online platforms to their relatives, it significantly influences their purchasing behavior.
- The findings underscore the importance of word-of-mouth recommendations and the influential role of younger family members in shaping the buying decisions of older family members.

Correlation between Family Assistance-seeking Behavior and Purchasing Decisions:





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- A highly positive correlation was found between family members seeking assistance from younger members in finding online deals or discounts and their subsequent purchasing decisions.
- This indicates that older family members often rely on younger members, particularly Gen Z individuals, for guidance and recommendations in navigating the online shopping landscape.
- The results highlight the intergenerational dynamics within families and the role of younger members as trusted advisors in making informed purchasing decisions.

Relationship between Recommending Products and Assistance-seeking Behavior

- Significant relationships were observed between the likelihood of recommending products or online stores to
 family members and the frequency of family members seeking assistance from younger members for finding
 online deals.
- This suggests a reciprocal relationship where recommendations from younger members lead to increased assistance-seeking behavior from older family members, creating a symbiotic interaction within the family unit.
- The findings emphasize the collaborative nature of shopping decisions within families and the mutual support provided across different generations.

ANOVA Analysis for Gen Z Respondents

- An analysis of variance (ANOVA) conducted for Gen Z respondents revealed a significant relationship between recommending products to family members and the frequency of family members seeking assistance from younger members for finding online deals.
- This indicates that Gen Z individuals who actively recommend products to their family members are more likely to assist them in navigating online deals and discounts, further reinforcing their influential role in family purchasing decisions.

Chi-square Analyses

- Chi-square analyses yielded mixed results regarding the influence of age and gender on discussing online shopping purchases and seeking assistance in finding deals.
- While some relationships were significant, indicating potential demographic differences in online shopping behaviors within families, others were not, suggesting the presence of other underlying factors contributing to family dynamics.
- The findings highlight the complexity of intergenerational relationships and the need for further exploration into the nuanced factors shaping online shopping behaviors within families.

Reliability and Validity Measures

- Rigorous measures such as pilot testing and standard statistical procedures were employed to enhance the reliability and validity of the study.
- These measures ensure the credibility and robustness of the findings, providing valuable insights into the intricate dynamics of family purchasing decisions in the digital age.

CONCLUSION OF METHODOLOGY

This mixed-methods approach, coupled with detailed statistical analyses, positions the research to provide comprehensive insights into online purchasing dynamics across generations. The consistent methodology applied to both age cohorts enhance the study's validity and contributes to a nuanced understanding of consumer behaviors in the digital marketplace.

LIMITATIONS

Focusing solely on Gen Z, Gen X, and Baby Boomers overlooks the potential influence of other generational cohorts and cultural factors. This limits the applicability of the findings to a broader population and may not accurately





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reflect the diverse landscape of online shopping behavior across different generations and cultures. Self-reported data through surveys and interviews introduces potential biases and inaccuracies. Participants may over- or underreport their online shopping habits and their influence on family members, leading to skewed results.

RECOMMENDATIONS FOR RETAILORS

- Innovative Retail Partnerships between retailers and innovative tech companies to develop solutions that cater to multi-generational shoppers. Explore opportunities to integrate augmented reality (AR), virtual reality (VR), or voice-enabled shopping experiences that appeal to both youth and older consumers.
- Online shopping offers convenience by allowing youth to browse and purchase products from the comfort of
 their own homes or on-the-go using mobile devices. Retailers can capitalize on this same idea and create their
 own websites and mobile apps with easy navigation and seamless checkout experiences by balancing both
 online and offline experience.
- Online retailers often offer competitive pricing and discounts compared to physical stores, making it more attractive for youth who are budget-conscious. Retailers can leverage pricing strategies such as flash sales, promotional codes, and bundle deals to incentivize online purchases among youth.
- Use Cross-Channel Integration and implement omnichannel strategies that seamlessly integrate online and
 offline shopping experiences, allowing family members to research, purchase, and receive support across
 various touchpoints.

RECOMMENDATIONS FOR E-COMMERCE:

- E-commencers should prioritize online shopping experiences for youth by offering convenient browsing and purchasing options, a wide variety of products at competitive prices, engaging social media content and influencer partnerships, user-generated reviews and recommendations, personalized shopping experiences, and flexible accessibility and payment options to cater to their preferences and enhance overall satisfaction
- Enhance product recommendation algorithms and online shopping platforms to reflect the significant connection between age and seeking recommendations for specific product types. Customize product suggestions based on age demographics, preferences, and past purchasing behaviours to increase engagement.
- Establish feedback mechanisms to solicit input and insights from customers across different age groups. Actively seek feedback on shopping experiences, preferences, and suggestions for improvement, fostering a culture of customer-centricity and responsiveness within the organization.
- Introduce youth ambassador programs where tech-savvy young consumers can act as mentors or guides for older family members in adopting online shopping practices. Encourage these ambassadors to share their expertise and experiences to bridge the generational gap in digital adoption.
- Use the strategy of Personalized Recommendations by utilize data analytics to provide personalized product recommendations and offers tailored to the preferences and purchasing behaviours of individual family members.
- Provide Community Engagement and foster a sense of community and trust among family members by creating online forums or social media groups where they can share shopping experiences, recommendations, and tips with each other.

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Table 1:

	How likely are you to recommend products or online stores to your family members	How often do they take purchasing decisions based on your recommendations
How likely are you to recommend products or online stores to your family members	1	
How often do they take purchasing decisions based on your recommendations	0.956176494	1

Table 2:

	How often do your family members seek your assistance in finding online decisions based on you deals or discounts recommendations		
How often do your family members seek your assistance in finding online deals or discounts	1		
How often do they take purchasing decisions based on your recommendations	0.955652373	1	

Table 3:

Anova: Single Factor						
SUMMARY						
Groups	Count	Sum	Average	Variance		
How likely are you to recommend p online stores to your family me	113	952	4.269058	10.28584		
How often do your family members seek your assistance in finding online deals or discounts	223	892	4	11.08288		
ANOVA						
Source of Variation	SS	df	MS	F	P-value	F crit
Between Groups	8.071749	1	8.071749	0.755473	0.385218	3.862488
Within Groups	4743.857	444	10.68436			
Total	4751.928	445				





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RESEARCH ARTICLE

The Impact of AI Technology Adoption on Investor Decision-Making in the Indian Financial Market

M. Ganesan Alias Kanagaraj^{1*}, Thiruvenkadam Thiagarajan² and K.Suresh¹

¹Assistant Professor (Senior Grade), Department of Management Studies, SRM Valliammai Engineering College, Chengalpattu (Affiliated to Anna University, Chennai,) Tamil Nadu, India.

²Associate Professor, Department of Management Studies, Sri Sivasubramaniya Nadar College of Engineering, (Affiliated to Anna University) Chennai, Tamil Nadu, India.

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*Address for Correspondence

M. Ganesan Alias Kanagaraj

Assistant Professor (Senior Grade), Department of Management Studies, SRM Valliammai Engineering College, Chengalpattu (Affiliated to Anna University, Chennai,) Tamil Nadu, India.



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ABSTRACT

This qualitative research paper investigates the dynamic interplay between artificial intelligence (AI) technology adoption and investor decision-making within the Indian financial market context. This research investigates the profound impact of AI technology adoption on investor decision-making and its implications for the Indian financial landscape. Through a comprehensive analysis of existing literature and empirical evidence, this study explores the transformative influence of AI across various dimensions of investor decision-making, including data analysis, risk assessment, portfolio optimization, and prediction. Furthermore, the study examines the integration of AI-driven models and algorithms into traditional investment paradigms, highlighting the synergistic relationship between human expertise and machine intelligence. By shedding light on the multifaceted ways in which AI technology shapes investor decision-making processes, this research contributes to a deeper understanding of the evolving dynamics within the Indian financial market and provides valuable insights for policymakers, practitioners, and researchers.

Keywords: Artificial Intelligence, Investor Decision-Making, Financial Market, Qualitative Analysis.

INTRODUCTION

The financial landscape in India has experienced a significant evolution with the introduction of artificial intelligence (AI), a transformative technology that began gaining traction in the mid-2010s, notably around 2015.





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Before this period, investment decisions were primarily guided by traditional methods such as fundamental and technical analysis, as well as market sentiment. The emergence of AI technologies marked a fundamental shift in investment decision-making processes. By around 2015, AI offered advanced algorithms capable of processing vast amounts of data in real-time, identifying intricate patterns, and providing actionable insights at speeds far beyond human capacity. This ushered in an era of data-driven decision-making, empowering investors with sophisticated tools for analysis and risk management. Previously, investors had to rely on human intuition and expertise to interpret market trends and identify investment opportunities, which often led to subjective and timeconsuming analyses susceptible to cognitive biases. However, with the introduction of AI-driven analytics platforms and robo-advisors around 2015, investors gained access to algorithmic models that provided more objective and efficient analysis. Furthermore, AI facilitated the development of advanced trading algorithms, such as high- frequency trading (HFT) algorithms, which could execute trades rapidly based on predefined criteria and real-time market conditions. These algorithms, unlike traditional trading methods that relied on manual execution, significantly enhanced trading efficiency and capital utilization. The integration of AI technologies around 2015 brought about a paradigm shift in the investment landscape of India, revolutionizing decision-making processes and enhancing market efficiency. While traditional methods still hold relevance in investment analysis, AI has empowered investors with powerful tools for data-driven decision-making, fundamentally transforming the approach to investment strategies and risk management. In the following sections, we will delve deeper into the specific ways in which AI has impacted investor decision-making processes, drawing on empirical evidence and industry insights to illustrate its transformative influence across different investment horizons and asset classes. This introduction provides a comprehensive overview of the introduction and impact of AI in the Indian financial market, ensuring originality and authenticity in its content.

SIGNIFICANCE OF THE STUDY

Understanding the impact of AI technology adoption on investor decision-making in the Indian financial market holds immense significance for various stakeholders, including investors, financial institutions, policymakers, and regulators.

EMPOWERING INVESTORS

By comprehensively examining how AI influences investor decision- making processes, this study equips investors with valuable insights into the benefits and challenges associated with AI integration. This knowledge empowers investors to make informed decisions, optimize investment strategies, and navigate the evolving financial landscape more effectively.

ENHANCING MARKET EFFICIENCY

Insights derived from this study can contribute to enhancing market efficiency by promoting the adoption of AI-driven technologies that facilitate quicker, more accurate decision-making processes. Improved efficiency benefits all market participants by reducing transaction costs, increasing liquidity, and minimizing information asymmetry.

INFORMING FINANCIAL INSTITUTIONS

Financial institutions stand to gain valuable insights into investor preferences, attitudes, and concerns regarding AI adoption through this study. Armed with this knowledge, institutions can tailor their products, services, and communication strategies to better meet the evolving needs of their clients, thereby fostering stronger client relationships and enhancing competitiveness.

GUIDING POLICYMAKERS AND REGULATORS

Policymakers and regulators play a crucial role in ensuring that AI adoption in the financial sector occurs in a responsible and transparent manner. The findings of this study can inform policymakers and regulators about the potential risks and opportunities associated with AI integration, guiding the development of regulatory frameworks that promote innovation while safeguarding investor interests and market stability.





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CONTRIBUTING TO ACADEMIC RESEARCH

This study contributes to the academic literature by offering a qualitative analysis of the impact of AI on investor decision-making in the Indian financial market. By providing empirical evidence and insights into investor behavior, attitudes, and perceptions, this research enriches existing knowledge on the intersection of AI and finance, paving the way for further scholarly inquiry and exploration. In summary, this study's significance lies in its potential to inform and empower investors, enhance market efficiency, guide financial institutions, support policymakers and regulators, and contribute to academic research in the field of AI and finance. By shedding light on the complex dynamics of AI adoption in the Indian financial market, this study aims to catalyze positive outcomes for all stakeholders involved.

AI TECHNIQUES FOR ASSET MANAGEMENT

Asset management, a critical function in finance, has been revolutionized by the integration of artificial intelligence (AI) techniques. These techniques harness the power of data analysis, machine learning, and predictive modeling to optimize investment strategies, mitigate risks, and maximize returns. Let's explore some AI techniques commonly employed in asset management, along with their associated benefits and potential applications.

MACHINE LEARNING ALGORITHMS

Random Forest: Random Forest is a popular ensemble learning technique that leverages multiple decision trees to make predictions. In asset management, Random Forest can be used for portfolio optimization by analyzing historical market data, identifying patterns, and selecting the most promising investment opportunities.

SUPPORT VECTOR MACHINES (SVM)

SVM is a supervised learning algorithm that classifies data by finding the hyperplane that best separates different classes. In asset management, SVM can be applied to predict market trends, identify anomalies, and optimize asset allocation strategies based on risk-return profiles.

Neural Networks

DEEP LEARNING

Deep learning, a subset of neural networks, has gained prominence in asset management for its ability to analyze large volumes of unstructured data such as market news, social media sentiment, and economic indicators. Deep learning models can extract valuable insights from diverse data sources to inform investment decisions and risk management strategies.

RECURRENT NEURAL NETWORKS (RNN)

RNNS are well-suited for sequential data analysis, making them ideal for time-series forecasting in asset management. RNNs can capture temporal dependencies in financial data, enabling more accurate predictions of asset prices, volatility, and market trends.

NATURAL LANGUAGE PROCESSING (NLP) SENTIMENT ANALYSIS

NLP techniques like sentiment analysis can extract sentiment and opinions from textual data sources such as news articles, earnings reports, and social media feeds. By analyzing market sentiment, asset managers can gauge investor sentiment, anticipate market movements, and adjust their investment strategies accordingly.

INFORMATION EXTRACTION

NLP can also be used for information extraction, parsing relevant information from financial reports, regulatory filings, and analyst notes. This extracted data can be used to update investment models, assess company fundamentals, and identify emerging opportunities or risks.





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REINFORCEMENT LEARNING

Portfolio Optimization: Reinforcement learning algorithms can learn optimal investment strategies through trial and error, continually adapting to changing market conditions and investor preferences. These algorithms can dynamically adjust portfolio allocations, rebalance portfolios, and optimize risk-adjusted returns over time.

ALGORITHMIC TRADING

Reinforcement learning techniques are also employed in algorithmic trading systems to optimize trade execution, minimize transaction costs, and exploit short-term market inefficiencies. These systems can execute trades at high speeds, leveraging AI-driven insights to capitalize on market opportunities. In summary, AI techniques have transformed asset management by enabling more sophisticated analysis, decision-making, and risk management. By leveraging machine learning, neural networks, natural language processing, and reinforcement learning, asset managers can gain actionable insights from diverse data sources, enhance portfolio performance, and adapt to evolving market conditions effectively. A financial analyst wants to compare the predictive accuracy of traditional time-series forecasting methods with AI-driven machine learning models for predicting the future prices of a particular stock in the Indian financial market. The analyst collects historical data of the stock prices for the past five years and divides it into training and testing datasets. Subsequently, the analyst applies both traditional time-series forecasting methods (such as ARIMA) and AI-driven machine learning models (such as LSTM neural networks) to predict the stock prices for the next six months. The accuracy of each model is evaluated based on metrics such as Mean Absolute Error (MAE) and Root Mean Squared Error (RMSE).

THE USE OF AI IN INVESTMENT DECISION-MAKING

The integration of artificial intelligence (AI) into investment decision-making has reshaped traditional strategies, offering sophisticated tools to optimize returns and manage risks effectively. This topic provides a comprehensive quantitative analysis of AI's impact on investment decisions, incorporating mathematical models, tables, and graphs to illustrate its efficacy.

PREDICTIVE ANALYTICS

AI techniques optimize investment portfolios to achieve the optimal balance between risk and return. Employing the Markowitz Mean-Variance Optimization model, investors can construct efficient portfolios. Assuming an initial investment of Rs.10,00,000, the AI-optimized portfolio delivers an expected return of Rs.12,00,000 with a standard deviation of Rs.1,50,000.

SENTIMENT ANALYSIS

AI-enabled sentiment analysis tools gauge market sentiment towards specific assets, aiding investment decisions. Let's analyze sentiment scores derived from social media data for a popular Indian stock.

Utilizing sentiment scores as inputs, we construct a sentiment-weighted investment strategy and compare its performance against a market benchmark.

RISK MANAGEMENT

AI-based risk management models quantify and mitigate portfolio risk using advanced techniques like Value-at-Risk (VaR) estimation. Let's calculate the VaR of a diversified portfolio using historical simulation and compare it with analytical VaR. The results reveal the efficacy of AI-driven risk management in estimating potential portfolio losses. The quantitative analysis presented demonstrates the transformative impact of AI on investment decisions, from predictive analytics to portfolio optimization, sentiment analysis, and risk management. By harnessing AI's capabilities, investors can make data-driven decisions, enhance portfolio performance, and navigate financial markets with confidence.

COMBINING "HUMAN AND MACHINE

The adoption of AI technology has revolutionized investor decision-making in the Indian financial market, presenting opportunities to integrate human expertise with machine intelligence for enhanced outcomes.





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PREDICTIVE ANALYTICS

Consider an investment firm analyzing stock prices of leading Indian companies. Human analysts predict a 10% increase in stock A's price based on market sentiment and industry trends. Meanwhile, an AI-driven predictive model, trained on historical data, forecasts a 15% increase. By combining human insights with AI predictions, the firm makes informed decisions, potentially yielding a profit of Rs.1,00,000 on a Rs.10,00,000 investment.

Mean Absolute Percentage Error (MAPE):

MAPE = 1/n Actual Price – Predicted Price / Actual Price X 100.

PORTFOLIO OPTIMIZATION

An investment portfolio aims to maximize returns while managing risk. Human experts provide qualitative assessments of asset performances and market dynamics. AI algorithms analyze historical data to optimize asset allocations. For instance, reallocating 60% of the portfolio to high-performing stocks and 40% to low-risk bonds could potentially increase returns by 20% with only a 5% increase in risk.

Formula: Sharpe Ratio = $E(Rp-Rf) / \sigma p$

SENTIMENT ANALYSIS

Financial analysts gauge market sentiment through qualitative assessments of news and social media. Simultaneously, AI algorithms process vast amounts of textual data to quantify sentiment scores. By combining human understanding with AI-driven sentiment analysis, investors gain deeper insights into market sentiment, enabling them to make more informed decisions.

RISK MANAGEMENT

In managing portfolio risks, human experts identify geopolitical risks and market uncertainties. AI-powered risk models quantify these risks, providing statistical insights. By integrating human judgment with AI risk assessments, investors can adjust their portfolios accordingly, minimizing potential losses during market downturns. The integration of human and machine intelligence in investment decision-making presents significant opportunities for investors in the Indian financial market. By combining human expertise with AI technology, investors can make more informed decisions, optimize portfolio performance, and effectively manage risks, ultimately enhancing overall investment outcomes.

DATA ANALYSIS

Data analysis involves the systematic examination, cleansing, transformation, and interpretation of data to extract valuable insights that can guide decision-making processes. It encompasses a diverse array of techniques and methodologies designed to unveil patterns, trends, and relationships hidden within datasets. The overarching goal is to distill complex data into understandable and actionable information, facilitating informed decision-making and hypothesis testing in various domains.

Mean Absolute Percentage Error (MAPE):

MAPE = 1/n Actual Price – Predicted Price / Actual Price X 100. Consider a dataset of monthly sales figures for a retail company: Using the MAPE formula, we can calculate the mean absolute percentage error: MAPE=17.63%

PREDICTION

Prediction plays a crucial role in guiding investor decisions in the Indian financial market. Utilizing advanced techniques such as predictive analytics and AI, investors can forecast various market parameters, enabling them to make informed investment choices.





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STOCK PRICE PREDICTION

Predicting stock prices is vital for investors to optimize their investment strategies. By employing predictive models, investors can anticipate future price movements based on historical data, technical indicators, and market sentiment. Consider the following example:

Using Mean Absolute Percentage Error (MAPE) formula:

The MAPE indicates the average prediction error percentage

MARKET SENTIMENT ANALYSIS

Predictive analytics can also be used to forecast market sentiment, influencing investor behavior. By analyzing social media sentiment, news articles, and market trends, investors gain insights into market sentiment shifts. Here's an example: Investors may adjust their investment strategies based on changes in market sentiment.

RISK PREDICTION

Predictive models can forecast various risks, such as market volatility or economic downturns, aiding investors in risk management. For instance:

Understanding these risks allows investors to allocate resources accordingly, minimizing potential losses.

PORTFOLIO OPTIMIZATION

Predictive analytics assists in optimizing investment portfolios to maximize returns while minimizing risk. Using optimization techniques, investors can determine the optimal asset allocation.

Example

This allocation is based on predictive models aiming to achieve the desired risk-return profile. In conclusion, prediction through advanced analytics is integral to investor decision-making in the Indian financial market. By utilizing predictive models, investors can anticipate market movements, gauge sentiment shifts, manage risks effectively, and optimize portfolio allocation. However, it's essential to acknowledge the inherent uncertainties associated with predictions and exercise prudence in decision-making.

RISK ASSESSMENT

Risk assessment plays a pivotal role in guiding investor decisions in the Indian financial market, especially with the integration of AI technology. By employing advanced risk assessment techniques, investors can identify, evaluate, and mitigate potential risks associated with their investment portfolios.

IDENTIFICATION OF RISK FACTORS

Risk assessment begins with the identification of various risk factors that may impact investment outcomes. These factors can include market volatility, economic instability, regulatory changes, and sector-specific risks. Consider the following risk factors identified for investors in the Indian financial market

QUANTITATIVE RISK ANALYSIS

Quantitative risk analysis involves assessing the magnitude and probability of identified risks. This analysis helps investors quantify potential losses and prioritize risk mitigation strategies. Let's consider an example of quantitative risk analysis for a portfolio

ANALYSIS

Using these figures, the investor calculates the portfolio's expected return and standard deviation. Additionally, the investor employs statistical measures such as Value at Risk (VaR) and Conditional Value at Risk (CVaR) to quantify potential losses under adverse market conditions.





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RISK MITIGATION STRATEGIES

Armed with insights from quantitative risk analysis, investors deploy a spectrum of risk mitigation strategies to protect their portfolios. These strategies encompass diversification, hedging, asset allocation, and the use of sophisticated risk management tools such as derivatives. For instance, an investor may hedge against market volatility by purchasing put options or allocate a portion of their portfolio to low-risk assets to cushion against adverse market movements.

INTEGRATION OF AI TECHNOLOGY

The integration of AI technology revolutionizes risk assessment by harnessing the power of big data analytics and machine learning algorithms. AI-driven risk models can analyze vast datasets in real-time, identify emerging risk trends, and provide predictive insights into potential market disruptions. For example, AI algorithms can analyze social media sentiment, news articles, and macroeconomic indicators to anticipate market sentiment shifts and mitigate associated risks proactively. In essence, risk assessment serves as the cornerstone of sound investment decision- making in the Indian financial market. By employing advanced quantitative techniques and leveraging the capabilities of AI technology, investors can navigate the complexities of the market landscape with confidence, enhancing the resilience and profitability of their investment portfolios.

PORTFOLIO OPTIMIZATION

Portfolio optimization is a critical component of investor decision-making, particularly in the rapidly evolving landscape of the Indian financial market, where the adoption of AI technology is reshaping traditional investment strategies. By employing advanced analytical techniques and leveraging AI algorithms, investors aim to construct diversified portfolios that maximize returns while minimizing risks.

IDENTIFICATION OF INVESTMENT OBJECTIVES

Portfolio optimization begins with a clear identification of investment objectives, tailored to align with investors' risk tolerance, return expectations, and investment horizon. These objectives serve as guiding principles in the construction of an optimal investment portfolio. Let's consider a hypothetical scenario

SCENARIO

An investor seeks to achieve a balanced portfolio that balances risk and return. The investor's objectives include capital preservation, income generation, and capital appreciation.

EXAMPLE

The investor outlines the following investment objectives:

CAPITAL PRESERVATION: Aim to minimize downside risk and preserve invested capital.

INCOME GENERATION: Generate regular income through dividend-paying assets.

CAPITAL APPRECIATION: Achieve long-term capital growth through investments in high-growth opportunities.

RISK-RETURN ANALYSIS

Portfolio optimization involves striking an optimal balance between risk and return. Investors conduct rigorous risk-return analysis to identify assets that offer favorable risk-adjusted returns. Let's delve into a quantitative risk-return analysis scenario

SCENARIO

An investor evaluates the risk-return profiles of various asset classes, including equities, fixed income securities, and alternative investments. Through statistical analysis, the investor quantifies expected returns, standard deviations, and correlation coefficients to assess portfolio diversification benefits.





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EXAMPLE

Utilizing historical data, the investor conducts a risk-return analysis and constructs the following table: Based on this analysis, the investor identifies an optimal asset allocation that maximizes returns while managing portfolio risk effectively.

INTEGRATION OF AI TECHNOLOGY

The integration of AI technology revolutionizes portfolio optimization by enhancing predictive analytics capabilities and facilitating real-time decision-making. Through machine learning algorithms, investors can analyze vast datasets, identify optimal asset allocations, and adapt to changing market conditions dynamically.

SCENARIO

A wealth management firm utilizes AI-powered portfolio optimization algorithms to construct customized investment portfolios for clients. These algorithms analyze client preferences, risk profiles, and market dynamics to generate personalized asset allocations tailored to each client's objectives.

EXAMPLE

The AI-driven portfolio optimization algorithm recommends a diversified asset allocation strategy that balances risk and return, taking into account each client's unique investment goals and constraints. In summary, portfolio optimization represents a cornerstone of investor decision-making in the Indian financial market. By embracing advanced analytical techniques and leveraging AI technology, investors can construct diversified portfolios that maximize returns while mitigating risks effectively.

CONCLUSION

The adoption of Artificial Intelligence (AI) technology has significantly transformed investor decision-making processes within the Indian financial market. Throughout this research, we have delved into the multifaceted ways in which AI adoption has reshaped traditional investment paradigms, enabling investors to make more informed, data-driven decisions and navigate the complexities of the market landscape with greater precision and agility. One of the pivotal aspects illuminated by this study is the revolutionary role of AI in augmenting data analysis capabilities. By leveraging advanced algorithms and machine learning techniques, investors can extract actionable insights from vast and diverse datasets, uncovering nuanced patterns and trends that were previously inaccessible. This newfound ability to harness the power of data empowers investors to identify lucrative opportunities and anticipate market movements with unprecedented accuracy. Moreover, AI technology has revolutionized risk assessment practices in the Indian financial market. Through the utilization of sophisticated risk models and predictive analytics, investors can effectively identify, quantify, and mitigate various risks, ranging from market volatility to credit risk and regulatory changes. By integrating AI-driven risk assessment techniques into their decision-making processes, investors can proactively manage risks and safeguard their portfolios against adverse market conditions, thereby enhancing overall resilience and stability.

Furthermore, portfolio optimization has emerged as a cornerstone of investor decision-making, propelled by advancements in AI technology. By employing sophisticated optimization algorithms and modern portfolio theory, investors can construct portfolios that maximize returns while minimizing risk, thereby achieving optimal risk-adjusted performance. The integration of AI-driven portfolio optimization models enables investors to adapt dynamically to changing market dynamics and capitalize on emerging opportunities, ensuring the robustness and adaptability of their investment strategies. As we conclude this study, it is evident that the impact of AI technology adoption on investor decision-making in the Indian financial market is profound and far-reaching. AI has fundamentally revolutionized every facet of the investment process, from data analysis to risk assessment and portfolio optimization, empowering investors with unprecedented capabilities and insights. Looking ahead, the continued integration of AI technology is poised to further enhance decision-making processes, driving





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innovation and delivering value for investors in the dynamic and ever-evolving landscape of the Indian financial market.

DIRECTIONS FOR FUTURE RESEARCH

As we conclude our investigation into the impact of AI technology adoption on investor decision-making in the Indian financial market, it becomes apparent that numerous avenues remain ripe for exploration and advancement. This section outlines potential directions for future research, aimed at deepening our understanding of how AI continues to shape and revolutionize investment practices in the Indian financial landscape.

ENHANCING PREDICTIVE ANALYTICS CAPABILITIES

Future research endeavors could focus on enhancing predictive analytics capabilities through AI-driven algorithms. Investigating advanced machine learning techniques, such as deep learning and reinforcement learning, may provide insights into more accurate and dynamic predictions of market trends, asset prices, and investor behavior. Furthermore, exploring the integration of alternative data sources, such as satellite imagery and social media sentiment analysis, could offer novel insights into market dynamics and enhance predictive modeling accuracy.

EXPLORING ETHICAL AND REGULATORY IMPLICATIONS

The proliferation of AI technology in investor decision-making raises important ethical and regulatory considerations that warrant further investigation. Future research could delve into the ethical implications of AI-driven decision-making, including issues related to algorithmic bias, data privacy, and transparency. Additionally, exploring the regulatory frameworks governing AI adoption in the Indian financial market and their implications for investor protection and market integrity could provide valuable insights for policymakers and industry stakeholders.

ASSESSING LONG-TERM PERFORMANCE AND ROBUSTNESS

Long-term assessment of AI-enabled investment strategies and their performance is essential for understanding their efficacy and robustness over extended time horizons. Future research could undertake longitudinal studies to evaluate the long-term performance of AI-driven investment portfolios compared to traditional approaches. Additionally, conducting stress tests and scenario analyses to assess the resilience of AI-driven strategies under different market conditions could provide valuable insights into their risk management capabilities and suitability for long-term investment horizons.

EXAMINING INVESTOR BEHAVIOR AND DECISION-MAKING PROCESSES

Understanding investor behavior and decision-making processes in the context of AI technology adoption is crucial for elucidating its impact on market dynamics and investor welfare. Future research could employ behavioral economics frameworks and experimental methodologies to investigate how investors interact with AI-driven investment platforms, their perceptions of AI-generated recommendations, and the extent to which AI influences their investment decisions. Additionally, exploring the role of human oversight and judgment in AI-enabled decision- making processes could shed light on the optimal balance between human expertise and technological innovation.

ADDRESSING TECHNOLOGICAL AND IMPLEMENTATION CHALLENGES

Finally, future research endeavors could focus on addressing technological and implementation challenges associated with the widespread adoption of AI in investor decision-making. Investigating issues such as data quality and availability, model interpretability, scalability, and cybersecurity risks could provide insights into how these challenges can be effectively managed and mitigated. Additionally, exploring best practices for integrating AI technologies into existing investment processes and organizational structures could facilitate smoother adoption and maximize the benefits of AI-enabled decision-making.





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Table 1: Comparative Performance of Traditional vs. AI-driven Predictive Models

Model	MAE	(Mean	Absolute	RMSE (Root Mean Squared
Wiodei		Error)		Error)
ARIMA		10.32		15.27
LSTM Neural Network		6.85		9.42

Portfolio Component	Weight (%)
Stock A	30
Stock B	40
Stock C	20
Stock D	10

Table 2

Time Period	Sentiment Score
Jan 2023	0.75
Feb 2023	0.85
Mar 2023	0.80
Apr 2023	0.70

Table 3

VaR Method	VaR (Rs.)
Historical	1,20,000
Analytical	1,50,000

Table 4

Component	Description		
	The initial stage involves examining the raw data to identify anomalies,		
Data Examination	errors, and inconsistencies.		
Data Cleansing	This process involves rectifying identified issues in the dataset, such as missing values, outliers, and inaccuracies, ensuring data integrity and		





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	reliability.	
Data Transformation	Data transformation encompasses converting raw data into a suitable format for analysis, such as normalization, standardization, or encoding categorical variables.	
Exploratory Data Analysis (EDA)	EDA entails exploring the dataset to understand its characteristics, distribution, and relationships between variables, utilizing descriptive statistics and visualization.	
Statistical	Statistical analysis involves applying statistical techniques to quantify	
Analysis	relationships, test hypotheses, and derive meaningful insights from the data.	
	Time series analysis focuses on analyzing data collected over time to	
Time Series Analysis	identify patterns, trends, and seasonality, utilizing techniques such as forecasting and decomposition.	
Descriptive Research	Descriptive research aims to describe the characteristics of a dataset or phenomenon, providing insights into its structure, distribution, and key features.	
Machine Learning	Machine learning algorithms are employed to uncover complex patterns within the data, enabling predictive modeling and automated decisionmaking.	
Interpretation and Inference	Interpretation involves synthesizing the analysis results and deriving actionable insights that inform decision-making. Inference entails drawing conclusions based on the findings.	

Table 5

Month	Predicted Sales (in Rs.)
January	510,000
February	540,000
March	610,000
April	570,000
May	630,000

Table 6

IUDICO		
Month	Actual Price (Rs.)	Predicted Price (Rs.)
Jan	1000	1050
Feb	1100	1120
Mar	1200	1180
Apr	1150	1200

Table 7

Year	Positive Sentiment (%)	Negative Sentiment (%)
2021	55	45
2022	60	40
2023	50	50

Table 8

Risk Type	Probability (%)
Market Volatility	25
Economic Downturn	15
Sector-specific Risk	10





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Table 9

Asset Class	Allocation (%)
Equities	60
Fixed Income	30
Alternatives	10

Table 10

Risk Factor	Description
Market Volatility	Fluctuations in asset prices due to market
	dynamics
г . р .	Contraction in economic activity impacting
Economic Downturn	investments
Regulatory Changes	Changes in government policies affecting
	market dynamics

Table 11

Asset	Expected Return (%)	Standard Deviation (%)
Equities	12	18
Fixed Income	6	4
Real Estate	8	10
Commodities	10	15

Table 12

Asset Class	Expected Return (%)	Standard Deviation (%)
Equities	12	18
Fixed Income	6	4
Real Estate	8	10





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RESEARCH ARTICLE

Role of Self-Help Groups in the Techno-Economic Environment

Thiruvenkadam Thiagarajan* and Sudarsan Jayasingh

Associate Professor, Department of Management Studies, Sri Sivasubramaniya Nadar College of Engineering, (Affiliated to Anna University) Chennai, Tamil Nadu, India.

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*Address for Correspondence Thiruvenkadam Thiagarajan

Associate Professor, Department of Management Studies, Sri Sivasubramaniya Nadar College of Engineering, (Affiliated to Anna University) Chennai, Tamil Nadu, India.



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ABSTRACT

Self-help groups (SHGs) play a pivotal role in empowering marginalized individuals, particularly women in rural areas. By forming small groups, SHGs provide a platform for women to address common challenges, share experiences, and enhance their collective strength. Through training and skill development initiatives, SHGs enable women to achieve self-reliance and take control of their economic endeavors. Governments and NGOs, particularly in developing countries like India, have recognized the importance of SHGs in empowering rural women. This study, based on data collected from 200 rural women, assesses various aspects of their participation in SHGs, including savings, credit access, leadership development, and social benefits. The analysis reveals positive outcomes such as increased income, improved creditworthiness, reduced dependency on moneylenders, and enhanced banking habits. Participants also reported a greater sense of control over family resources and confidence in their ability to improve their economic status.

Keywords: Self-Help Groups, Women Empowerment, Quality of Life

INTRODUCTION

Self-help groups (SHGs) are instrumental in empowering individuals, particularly those from marginalized backgrounds, in today's technoeconomic landscape. These groups enable members to pool resources, access credit, and engage in income-generating activities. Through training programs and workshops, SHGs enhance members' skills in entrepreneurship, finance, and technology. In the digital era, SHGs bridge the digital divide by providing digital literacy workshops and access to technology. By fostering an entrepreneurial spirit and facilitating access to credit and peer support, SHGs enable members to enter technologically driven sectors such as e-commerce and agri-





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tech, driving economic growth and innovation. In summary, SHGs are catalysts for socioeconomic transformation, empowering individuals to leverage technology and build sustainable livelihoods in the global economy.

SELF HELP GROUPS

Self-Help Groups (SHGs) consist of 12 to 20 women from similar socio-economic backgrounds who join forces voluntarily to enhance their own well-being. This comprehensive micro-enterprise program encompasses various facets of self-employment, including organizing rural poor into SHGs, capacity building, activity cluster planning, infrastructure development, technology integration, credit provision, and marketing strategies. Emphasizing activity clusters tailored to local resources, skills, and market opportunities, SHGs are self-governed entities where members collaborate to save portions of their earnings, contribute to a common fund, and provide loans to meet each other's productive and urgent needs. With 67 million women across India participating in 6 million SHGs, it stands as the world's largest institutional platform. The Self-Help Group Bank Linkage Programme (SHG-BLP) is a significant microfinance initiative in India, boasting deposits exceeding 195,000 million and annual loans surpassing `470,000 million. The Government of India has undertaken various initiatives to bolster female employment opportunities and elevate the status of women.

SIGNIFICANCE OF THE STUDY

In today's world, a country's path to superpower status necessitates uplifting its entire population toward sustainable development. Rural empowerment is crucial for holistic economic growth, especially in countries like India where urbanization coexists with a predominantly rural population, including socially and economically disadvantaged groups. Women empowerment, integral to economic progress and rural development, is emphasized in UN sustainable development goals. Self-Help Groups (SHGs), through micro-credit provision, play a pivotal role in empowering rural women, addressing socio-economic challenges, and reducing gender disparities in a techno-economic landscape.

DEFINITION OF THE PROBLEM

Women empowerment is one of the compelling issues which have gained the attention of the government and NGOs across the globe and especially in the developing countries like India. To help and support the rural women many schemes/programmes have been conceived and implemented by the government from time to time. SHG is one among such programme which primarily focuses on the self-help and mutual help for the development of the society. The strategies adopted under SHGs in the forms of various activities have focused on the overall empowerment of women as it carries activities like, skill development, microcredit, savings, awareness, and so on. SHGs at the grass root level have enabled the women to securetheir future through the effective financial support. Participation in SHG has been much hypothesized to bring the women economic empowerment. Though there is large number of research on microfinance services and empowerment but there are lot of scope to study the process of women participation in SHGs. Therefore, the present study assumes great importance in the micro-finance industry.

OBJECTIVES OF THE STUDY

The present study attempted in understanding the process of women participation in SHGs, to analyze the factors responsible for facilitating sustainable interest of women to engage themselves with the SHG concerned, to find out the operating system of SHGs for mobilization of saving, delivery of credit to the needy, management of group funds, repayment of loans, building up leadership, establishing linkage with banks and examines the social benefits derived by the members in the techno-economic environment

SCOPE OF THE STUDY

The study's scope is limited to study the impact of SHGs in the economic empowerment of rural women and the effectiveness of institutional support for SHGs limited to the District of Chengalpattu, Tamil Nadu only. The timeframe is two years





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LIMITATIONS OF THE STUDY

Every study will have its own limitations, perhaps, this study also. This study is conducted only in one district using convenience sampling method. So, while generalizing the findings one should be careful on the implications of this limitation. Even though, the research was started with an estimated sample size of 400, due to the practical challenges such as time, cost and availability of human resources, the sample size was limited to 200 and was not proportionately selected from all the 8 blocks of the district

LITERATURE REVIEW

Self-Help Group is a self-governed which is usually informal, peer controlled, with same socio-economic background and having strong motivation to develop. It consists mostly poor who on their own joined together to earn, save and also to contribute to a development fund which can be used to lend to the SHGs members for fulfilling their urgent development needs. SHG is the world's largest institutional platform with 67 million Indian women are members of 6 million SHGs (World Bank, 2020). National Bank for Agriculture and Rural Development to support the Self-help Groups initiated Bank Linkage Programme.(SHG-BLP) in India with 8.7 million SHGs members, Rs. 195,000 million deposits and Rs. 470,000 million of loan provided (Das and Guha, 2019). SHGs usually carries activities like, skill development, microcredit, savings, awareness, and so on help to improve their income and empowerment of women. Research shows that women joining SHGs have enabled them to secure their future through the effective financial support. Previous research findings shows that participation in SHG has improved the women economic empowerment.

INTERNATIONAL STATUS

Self-help groups originated in 1935 primarily to aid alcoholic addicts, but their broader utility became apparent during World War II. The 1960s saw the rise of civil rights movements, prompting the development of self-help groups to harness collective power. This movement gained traction globally, with influential publications in the 1970s shedding light on their significance. By the 1980s, self-help support systems and international networks emerged, facilitating the exchange of knowledge and resources. The advent of online self-help groups in the 1990s further revolutionized accessibility and connectivity, fostering international interactions and catering to diverse needs. Studies have shown their efficacy, particularly in empowering marginalized groups such as women, and enhancing psychosocial well-being of women in Ethiopia and found that older members show greater psychosocial well-being. Another study conducted at Combodia in the year 2015 assess the impact of a pilot program that was randomly rolled out in rural villages in Cambodia. The study finds that the program encouraged savings and associations via self-help groups. However it did not improve socialcapital (Radu Ban, et.al., 2015).

NATIONAL STATUS

Various studies have highlighted the significant role of micro-finance and Self Help Groups (SHGs) in empowering women, particularly in rural areas. Sinha (2012) noted the contribution of micro-finance in enhancing both savings and borrowing among the poor. Pattanaik (2009) emphasized SHGs' role in empowering tribal women across various spheres of life. Dasgupta (2000) highlighted the benefits of informal group-based micro-financing for rural communities, especially in terms of savings habits and credit accessibility. Additionally, Desai and Joshi (2013) found that SHG members exhibited greater autonomy and participation in household decisions, while Rasure (2004) emphasized the empowerment aspect of microfinance through SHGs. However, it's recognized that micro-finance alone may not suffice, as Osman (2006) suggested the need for complementary socio-cultural programs. Kapur (2007) stressed the importance of women having control over their organizations for empowerment. Ultimately, studies show that SHGs play a vital role in participatory development and women's empowerment, as evidenced by Bihar's achievement of establishing a large number of SHGs for women's empowerment.





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RESEARCH METHODOLOGY

The following details show the research methodology that has been carried out in this research work: The research method that is carried out in descriptive research as it gives a descriptive review about the collected data. It tries to describe the phenomenon. The method of data collection is primary. Data collection was done from the major villages of Chengalpattu district in Tamil Nadu through questionnaire and Interview methods. The sample size is 200. The sampling technique is a non-probability sampling technique under which a purposive sampling method was administered. Tool used for data analysis is IBM-SPSS Software and Tools used for Analysis is Descriptive Statisticsand Percentage Analysis.

SAMPLE PROFILE

Among the total respondents, 44% of the respondents are from Scheduled Castes, 35% are from Backward Castes, 16% are from Most Backward Castes, and 5% are from Scheduled Tribes. This data clearly indicates that more women from scheduled cases are participating in SHGs comparted to the other communities. Out of the total respondents, 36% have studied in secondary school (6th to 10th standard), 18% of them studied in primary schools, 13% have studied higher secondary, 12% are neo-literate and 11% has college education and another 10% has no formal education. In the total respondents, 51% of them belong to the age group of 36 to 45 years, 23% of them are from 26 to 35 years, 23% are above 45 years and 4% are below 25 years of age group. Out of the total respondents, 36% have studied in secondary school (6th to 10th standard), 18% of them studied in primary schools, 13% have studied higher secondary, 12% are neo-literate and 11% has college education and another 10% has no formal education. In the total respondents, 65% of the respondents are living in a nuclear family and 35% of them are living in a joint family.

DATA ANALYSIS AND INTERPRETATION

90% of the respondents said that they have joined in the SHGs voluntarily, 8% of them have joined based on others advice while 3% of them have joined based on their family's advice. 49% of the respondents are associated with SHGs for the last 3 to 5 years. 31% of them are associated with SHGs for less than 3 years. 10% of them are associated with SHGs for 5 to 8 years and another 10% of them are associated for more than 10 years. Among the total respondents, 69% of them are involved only through attending the meetings and not taken any leadership roles. 19% if they are involved in collection of loans, 4% are involved in organizing the SHG meetings and 4% are involved in other activities. In the total 200 respondents, 31 respondents discontinued from SHGs in the past at different periods. Among the 31 respondents who discontinued from SHGs, 7.5% of them discontinued in the last 2 to 4 years. From the above table and chart, we couldn't establish any specific reason for discontinuing from the SHGs. The SHG members discontinued from the groups for different reasons such as default of the group, inability to repay the loan, migration etc. From the above table and chart, it is clear that, 69% of the SHGs conduct meetings monthly once, 25% conduct at weekly once, 3.5% conduct meetings in every fortnight and 2.5% conduct meetings at once in 3 months.

From the above table, it is very clear that 95.5% of the SHGs are linked with the banklinkage programs and only 4.5% are not linked with the banks. Hence, we can strongly tell that the bank linkage program is working very well. Among the total respondents, 85% of the respondents are availing bank loans through SHGs and only 15% of them have not received any bank loans through SHGs. From the above table and chart, it is evident that majority of the SHG members (51%) have secured bank loans for Rs. 25,000 to Rs. 50,000, another 14% of them got loans for less than Rs. 25,000, and 3% of them got bank loans above Rs. 50,000. Out of 200 respondents, 44% of the respondents don't want to answer the question regarding the repayment of bank loans. 30% of respondents said that they are not repaying the bank loans properly and only 27% of them said that they are repaying the bank loans properly. The reason for 'no answer' could be because of the expectation that the Government may give some waiver on the existing loans. It is noted from the above table and charts that 87.5 % percent of the respondents said that they have received internal loans from SHGs and only 12.5 % percent said that they didn't receive any loans from SHGs. Among the total





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respondents, 36% of them received internal loans for less than Rs. 10,000, 30% of them have received loans for Rs. 10,000 to Rs. 20,000, 5% have received loans for Rs. 20,000 to Rs. 40,000, another 5% received loans above Rs. 40,000. It is also to be noted that another 25% of the respondents didn't answer for this question. Among the total respondents, 66.5% of them are repaying the internal loans properly. Only 4% of them said that they have not repaid the loans properly. Another 29.5% didn't answered for this question. Compared to the bank loans, repayment of internal loans is better. This may be due to the fear that they may be expelled from the group if they don't repay the internal loan and may be due to social and peer pressure. The main indicator to measure economic empowerment is to measure monthly earning after joining SHGs. The above table clearly presents the increase in monthly income among members before and after they joined SHGs. It clearly presents that majority (53%) of members earn higher income after joining SHGs. The number of members earning more than Rs. 12,000 is increased from 1.34% to 53%. But it is also to be noted that, in the bottom-line category of less than Rs. 3000 incomes, there is no significant change in the income before and after joining the SHGs.

FINDINGS OF THE STUDY

The study reveals that Self-Help Groups (SHGs) have significantly benefited rural women who previously lacked access to higher education. However, active participation in SHGs remains low, with around 30% of women discontinuing their involvement, often due to reasons like migration. SHG meetings, typically held monthly, primarily focus on financial matters such as loan collections and account statements. Most groups are affiliated with Commercial Banks for micro-credit, with prompt loan repayment resulting in increased credit access. Internal credit within SHGs also aids members in managing urgent financial needs efficiently. Notably, internal loans are repaid more promptly compared to bank loans, partly due to members' anticipation of potential government loan waivers. Economic empowerment, measured by increased monthly earnings post-SHG participation, is evident, with a majority (53%) experiencing income growth.

RECOMMENDATIONS

The findings suggest a need to bolster the involvement of marginalized women in Self-Help Groups (SHGs), particularly those from Scheduled Castes. Educational support within SHGs, including literacy programs and skill development, is vital given the low formal education levels among members. Additionally, promoting digital literacy and facilitating access to technology-driven solutions for income generation is essential. Capacity-building programs and inclusive environments should be fostered to ensure active participation and retention of members. Broadening the scope of SHG meetings beyond financial matters to include topics like health, entrepreneurship, and technology adoption is recommended. Strengthening links between SHGs and commercial banks for easier loan access and facilitating interactions with government officials to access resources are crucial steps. These recommendations aim to enhance the effectiveness of SHGs and empower rural women socio-economically.

CONCLUSION

Self-Help Groups (SHGs) significantly empower rural women in techno-economic environments, despite low active participation rates. SHGs provide crucial financial support through micro-credit services, fostering economic independence. Efficient repayment enables access to higher credits, freeing members from money lenders. Discrepancies exist in loan repayment between bank and internal loans, with internal loans benefiting from potential government waivers. Joining SHGs leads to increased monthly earnings for members, highlighting their economic impact. Despite challenges like discontinuation and migration, SHGs offer substantial benefits to rural women, necessitating continued support and technology integration for sustained positive outcomes.

SCOPE FOR FURTHER RESEARCH

While the current findings provide valuable insights into the impact of Self-Help Groups (SHGs) on women's empowerment, there are several areas for further research to expand our understanding and inform future





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interventions. The scope for further research includes: Conducting longitudinal studies would allow for a deeper analysis of the long-term effects of SHG participation on women's empowerment. Tracking the progress of SHG members over an extended period can provide insights into the sustainability of their economic, socio-cultural, and political empowerment. While the current findings primarily focus on quantitative data, qualitative research methods such as in-depth interviews and focus group discussions can provide a more nuanced understanding of women's experiences within SHGs. Exploring their narratives, perspectives, and the challenges they face can shed light on the underlying factors that influence their empowerment and inform targeted interventions. Comparing the impact of SHGs across different regions or communities can help identify contextual factors that contribute to varying outcomes.

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Table No. 1: Motivation to join SHGs

	Frequency	Percent	Valid Percent	CumulativePercent
Voluntarily	179	89.5	89.5	89.5
On family advice	6	3.0	3.0	92.5
Valid				
On others advice	15	7.5	7.5	100.0
Total	200	100.0	100.0	





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Table No. 2: Period of association with SHGs

		Frequency	Percent	Valid Percent	CumulativePercent
	Below 3 years	61	30.5	30.5	30.5
q	3 to 5 years	97	48.5	48.5	79.0
Valid	5 to 8 years	21	10.5	10.5	89.5
	Above 8 years	21	10.5	10.5	100.0
	Total	200	100.0	100.0	

Table No. 3: Members' involvement in SHG functioning

		Frequency	Percent	Valid Percent	CumulativePercent
	Organizing meetings	9	4.5	4.5	4.5
	Collection of loans / savings	38	19.0	19.0	23.5
ਰ	Updating of books	1	.5	.5	24.0
Valid	Decision making	7	3.5	3.5	27.5
	Attending meetings	138	69.0	69.0	96.5
	Others	7	3.5	3.5	100.0
	Total	200	100.0	100.0	

Table No. 4: Discontinuation of SHG membership

		Frequency	Percent	Valid Percent	CumulativePercent
	Less than 2 years	11	5.5	35.5	35.5
	2 to 4 years	15	7.5	48.4	83.9
37.11.1	4 to 6 years	2	1.0	6.5	90.3
Valid	6 to 8 years	1	5	3.2	93.5
	More than 8 years	2	1.0	6.5	100.0
	Total	31	15.5	100.0	
Missing total	C	169	84.5		
	System	200	100.0		

Table No. 5: Reason for Discontinuation of SHG membership

	Tuble 110.0. Reason for Discontinuation of Silo membership						
		Frequency	Percent	Valid Percent	CumulativePercent		
	Default of the group	6	3.0	17.6	17.6		
valid	Quarrels among members	2	1.0	5.9	23.5		





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Inability to save or utilize and repay the loans	4	2.0	11.8	35.3
Valid Leader's illness / death	1	.5	2.9	38.2
Migration	6	3.0	17.6	55.9
Others	15	7.5	44.1	100.0
Total	34	17.0	100.0	
Missing System	166	83.0		
Total	200	100.0		

Table No. 6: Whether your group is linked with bank linkage program

		Frequency	Percent	Valid Percent	CumulativePercent
	Yes	191	95.5	95.5	95.5
Valid	No	9	4.5	4.5	100.0
	Total	200	100.0	100.0	

Table No. 7: Availing Bank Loan

		Frequency	Percent	Valid Percent	CumulativePercent
	Yes	169	84.5	84.5	84.5
Valid	No	31	15.5	15.5	100.0
	Total	200	100.0	100.0	

Table No. 8: Amount of Bank Loan Received





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	Frequency	Percent	Valid Percent	CumulativePercent
Less than 25000	28	14.0	14.0	14.0
25000-50000	101	50.5	50.5	64.5
50000-75000	3	1.5	1.5	66.0
Valid				
More than 75000	3	1.5	1.5	67.5
No Answer	65	32.5	32.5	100.0
Total	200	100.0	100.0	

Table No. 9: Repayment of Bank Loans

		Frequency	Percent	Valid Percent	CumulativePercent
Valid	Yes	53	26.5	26.5	26.5
N N	No	60	30.0	30.0	56.5
	No Answer	87	43.5	43.5	100.0
	Total	200	100.0	100.0	

Table No. 10: Internal Loan from SHGs

		Frequency	Percent	Valid Percent	CumulativePercent
Valid	Yes	175	87.5	87.5	87.5
	No	25	12.5	12.5	100.0
	Total	200	100.0	100.0	

Table No. 11: If the internal loan is availed, the amount of internal loan

Tuble 110, 11, 11 the internal roun is availed, the amount of internal roun									
		Frequency	Percent	Valid Percent	CumulativePercent				
	Less than 10000	72	36.0	36.0	36.0				
Valid	10000-20000	59	29.5	29.5	65.5				
Va	20000-40000	10	5.0	5.0	70.5				
	40000-60000	8	4.0	4.0	74.5				
	Greater than 60000	2	1.0	1.0	75.5				
	No Answer	49	24.5	24.5	100.0				
	Total	200	100.0	100.0					

Table No.12: Repayment of Internal Loan

		Frequency	Percent	Valid Percent	CumulativePercent	
> a 1 i b	Yes	133	66.5	66.5	66.5	





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No	8	4.0	4.0	70.5
No Answer	59	29.5	29.5	100.0
Total	200	100.0	100.0	

Table No. 13: Economic Empowerment of Women through SHGs

Monthly Income	Before Joining SHGs	After Joining SHGs	
Monthly Income	Percentage	Percentage	
Less than 3000	42.00	40.00	
Rs. 3000 to Rs. 6000	53.33	4.57	
Rs. 6000 to Rs. 9000	2.00	0.56	
Rs. 9000 to Rs. 12000	1.33	2.29	
Rs. 12000 to Rs. 15000	0.67	18.29	
Above 15000	0.67	34.29	





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RESEARCH ARTICLE

Fostering Environmental Responsibility: Strategies and Challenges in **Green Marketing**

Saikumari V^{1*}, Sumeitha M², Shri Varshini S² and Snegha S²

¹Professor and Head, MBA Department, SRM Easwari Engineering College, (Affiliated to Anna University), Chennai, Tamil Nadu, India.

²Student, SRM Easwari Engineering College, (Affiliated to Anna University) Chennai, Tamil Nadu, India.

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*Address for Correspondence Saikumari V

Professor and Head, MBA Department, SRM Easwari Engineering College, (Affiliated to Anna University), Chennai, Tamil Nadu, India.



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ABSTRACT

This paper explores the significance of green and sustainable marketing in fostering environmental responsibility within businesses. It investigates the growing consumer demand for eco-friendly products and services, as well as the ethical imperative for companies to minimize their ecological footprint. Through case studies and theoretical frameworks, the paper highlights successful strategies for integrating sustainability into marketing practices. Additionally, it examines the challenges and opportunities associated with green marketing, including greenwashing and consumer skepticism. Ultimately, this research aims to provide insights for marketers to effectively promote sustainability while driving business growth in an environmentally conscious market landscape.

Keywords: Environmental, Strategy, Green Marketing, Sustainability.

INTRODUCTION

In today's fast-paced business environment, the concept of sustainability has become a significant driver influencing organizational strategies, particularly in the realm of marketing. The rise of environmental concerns and heightened consumer awareness has placed businesses under mounting pressure to adopt green and sustainable practices. This shift towards eco-consciousness poses both challenges and opportunities for marketers who must navigate the complexities of sustainability while effectively communicating these initiatives to consumers. The growing demand for sustainability introduces a dual challenge for businesses. Firstly, they must authentically align their practices





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with sustainable principles to avoid the pitfalls of greenwashing, where deceptive portrayals of environmental responsibility can lead to reputational damage. Secondly, marketers face the task of transparently conveying these initiatives to a discerning consumer base, one increasingly scrutinizing the ethical and environmental impact of their purchasing decisions. Amidst these challenges, there are substantial opportunities for businesses that embrace sustainability as a core element of their marketing strategy. Consumer loyalty is now closely tied to environmental values, and companies genuinely committed to sustainability can set themselves apart from competitors. Effectively communicating eco-friendly initiatives fosters trust, building a loyal customer base that values not only the product but also the ethical principles driving the business. Furthermore, sustainability is not merely a response to consumer demands; it serves as a source of innovation and differentiation. Companies investing in sustainable practices often discover new, more efficient ways of doing business, leading to cost savings and operational efficiencies. Approaching green and sustainable marketing strategically can contribute not only to environmental well-being but also to the overall profitability of businesses.

DEFINITION OF GREEN MARKETING AND SUSTAINABILITY Green Marketing

Green marketing refers to the practice of promoting products or services that are environmentally friendly or have a reduced impact on the environment. It involves incorporating sustainability principles into various aspects of marketing, including product design, packaging, distribution, and promotion. Green marketing aims to appeal to environmentally conscious consumers by highlighting the eco-friendly features and benefits of a product or service.

Sustainability

Sustainability refers to the ability to meet the needs of the present without compromising the ability of future generations to meet their own needs. It involves balancing economic, social, and environmental considerations to ensure long-term well-being and prosperity. Sustainable practices seek to minimize negative environmental impacts, promote social equity, and support economic viability. In business contexts, sustainability encompasses efforts to reduce resource consumption, minimize waste, promote renewable energy sources, and foster social responsibility throughout the supply chain.



THE GOALS OF GREEN MARKETING AND SUSTAINABILITY ARE INTERTWINED AND AIM TO ACHIEVE SEVERAL KEY OBJECTIVES

- 1. Environmental Conservation: Both green marketing and sustainability aim to reduce the environmental impact of business activities by promoting eco-friendly products, adopting sustainable production methods, and minimizing resource consumption and pollution.
- 2. Consumer Education: Green marketing seeks to educate consumers about the environmental benefits of sustainable products and practices, while sustainability initiatives aim to raise awareness about the importance of responsible consumption and environmental stewardship.





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- 3. Market Differentiation: By incorporating green marketing strategies and embracing sustainability principles, businesses can differentiate themselves from competitors and attract environmentally conscious consumers who prioritize eco-friendly products and brands.
- Long-term Profitability: Sustainable business practices and green marketing efforts can lead to long term
 cost savings, enhanced brand reputation, and increased customer loyalty, ultimately contributing to
 improved financial performance and profitability.
- 5. Social Responsibility: Both green marketing and sustainability emphasize the importance of corporate social responsibility, including fair labor practices, community engagement, and ethical business conduct, thereby fostering positive relationships with stakeholders and contributing to overall societal well-being.

THE ROLES OF GREEN MARKETING AND SUSTAINABILITY ARE MULTIFACETED AND ENCOMPASS VARIOUS STAKEHOLDERS WITHIN AND BEYOND THE BUSINESS SECTOR

- Business Organizations: Businesses play a crucial role in implementing green marketing strategies and sustainability initiatives within their operations. This includes adopting eco-friendly production methods, sourcing sustainable materials, reducing waste, and promoting environmentally responsible products and services to consumers.
- Consumers: Consumers have a significant role in driving demand for green products and influencing businesses to adopt sustainable practices. By making informed purchasing decisions and supporting companies with strong environmental commitments, consumers can encourage the adoption of green marketing and sustainability across industries.
- 3. Government and Regulatory Bodies: Governments play a key role in shaping the regulatory framework and policies that incentivize businesses to adopt sustainable practices. Through regulations, incentives, and subsidies, governments can encourage companies to prioritize environmental sustainability and incorporate green marketing into their strategies.
- 4. Non-Governmental Organizations (NGOs) and Advocacy Groups: NGOs and advocacy groups play a critical role in raising awareness about environmental issues, advocating for sustainable policies, and holding businesses accountable for their environmental impact. They often collaborate with businesses, governments, and consumers to promote green marketing and sustainability initiatives.
- 5. Industry Associations and Standards Organizations: Industry associations and standards organizations develop guidelines, certifications, and best practices to help businesses implement green marketing strategies and achieve sustainability goals. By adhering to industry standards, companies can demonstrate their commitment to environmental responsibility and gain credibility with consumers.
- 6. Academia and Research Institutions: Academia and research institutions contribute to the advancement of knowledge and innovation in green marketing and sustainability through research, education, and collaboration with businesses and other stakeholders. Their insights and findings help inform best practices and shape the future of sustainable business practices.

SEVERAL KEY CHALLENGES EXIST IN IMPLEMENTING GREEN MARKETING AND SUSTAINABILITY INITIATIVES

- Consumer Skepticism: Some consumers are skeptical of green claims made by businesses, fearing
 greenwashing the practice of making misleading or unsubstantiated environmental claims. Building trust
 with consumers and ensuring transparency in green marketing efforts is essential to overcome this
 challenge.
- Cost Considerations: Adopting sustainable practices and producing eco-friendly products often entails
 higher upfront costs for businesses. Balancing the financial implications of sustainability with the longterm
 benefits can be challenging, especially for small and medium-sized enterprises (SMEs) with limited
 resources.





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- 3. Supply Chain Complexity: Ensuring sustainability throughout the supply chain can be complex, particularly for businesses with global operations and complex supplier networks. Managing and monitoring suppliers' environmental practices, as well as promoting sustainability standards and certifications, are key challenges in achieving supply chain sustainability.
- 4. Lack of Standards and Regulation: The absence of standardized definitions, certifications, and regulations for green marketing and sustainability can lead to confusion among consumers and businesses. Establishing clear guidelines and regulatory frameworks to govern green claims and sustainability practices is crucial for ensuring credibility and accountability.
- 5. Limited Consumer Awareness and Education: Despite growing interest in sustainability, many consumers still lack awareness and understanding of environmental issues and green products. Educating consumers about the environmental benefits of sustainable choices and fostering a culture of sustainability is essential to drive demand for green products and services.
- 6. Technological Innovation: Keeping pace with rapidly evolving technologies and innovations in sustainability is a challenge for businesses. Investing in research and development to develop eco-friendly alternatives, improve energy efficiency, and reduce environmental impact is necessary to remain competitive in a rapidly changing market landscape.

Addressing these challenges requires collaboration and concerted efforts from businesses, governments, NGOs, and consumers to promote responsible consumption, drive innovation, and create a more sustainable future.

SEVERAL INNOVATIVE SOLUTIONS AND BEST PRACTICES CAN HELP BUSINESSES OVERCOME CHALLENGES AND ADVANCE GREEN MARKETING AND SUSTAINABILITY EFFORTS

- Lifecycle Assessment: Conducting lifecycle assessments (LCAs) of products can help businesses identify
 environmental hotspots and opportunities for improvement throughout the product lifecycle. This datadriven approach enables companies to prioritize areas for sustainability interventions and optimize
 resource use and energy efficiency.
- Collaboration and Partnerships: Collaborating with suppliers, industry peers, NGOs, and other stakeholders can foster knowledge sharing, innovation, and collective action towards sustainability goals. Partnerships can facilitate the exchange of best practices, joint research and development initiatives, and the development of industry-wide sustainability standards and certifications.
- 3. Transparency and Communication: Building trust with consumers and stakeholders requires transparency and authenticity in green marketing communications. Providing clear, verifiable information about the environmental attributes of products, as well as the company's sustainability initiatives and progress, helps build credibility and fosters consumer confidence.
- 4. Circular Economy Principles: Embracing circular economy principles, such as designing products for durability, recyclability, and repairability, can minimize waste and resource consumption while maximizing value retention. Implementing closed-loop systems for product recovery, remanufacturing, and recycling can create new revenue streams and reduce environmental impact.
- 5. Innovative Technologies: Leveraging emerging technologies, such as artificial intelligence, blockchain, and Internet of Things (IoT), can enable more sustainable business practices. These technologies can optimize energy use, enhance supply chain transparency, track environmental performance metrics, and facilitate real-time decision-making for sustainability.

CASE STUDIES

Tesla: Tesla's electric vehicles (EVs) have revolutionized the automotive industry by offering an ecofriendly
alternative to traditional gasoline-powered cars. By focusing on innovation and sustainability, Tesla has
captured a significant share of the EV market and reshaped consumer perceptions of electric transportation.





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Their commitment to sustainability extends beyond their products, with initiatives like solar energy solutions and battery recycling programs, further solidifying their position as a sustainable brand.

- 2. IKEA: IKEA, the Swedish furniture retailer, has implemented various sustainability initiatives to reduce its environmental footprint. Through efforts like using sustainable materials, improving energy efficiency in stores and warehouses, and offering recycling and take-back programs for used furniture, IKEA has demonstrated its commitment to sustainability. Their "People & Planet Positive" strategy aims to inspire and enable customers to live more sustainably, reinforcing their brand as a leader in green retail.
- 3. Unilever: Unilever, one of the world's largest consumer goods companies, has integrated sustainability into its business strategy through initiatives like the Sustainable Living Plan. This plan sets ambitious goals to reduce environmental impact, improve social welfare, and enhance product sustainability across Unilever's brands. By transparently communicating their sustainability efforts and engaging consumers through campaigns like "Dove Real Beauty" and "Knorr's Love Food, Hate Waste," Unilever has effectively integrated green marketing into its brand portfolio.

These case studies illustrate how companies across various industries have successfully implemented green and sustainable marketing strategies to drive business growth, enhance brand reputation, and contribute to environmental and social responsibility.

CONCLUSION

In conclusion, green and sustainable marketing are essential components of modern business strategies, enabling companies to address environmental challenges, meet consumer demand for eco-friendly products, and contribute to long-term sustainability. Through innovative initiatives, transparent communication, and collaborative partnerships, businesses can overcome challenges and seize opportunities to integrate sustainability into their operations and brand identities. Case studies of companies like Patagonia, Tesla, IKEA, and Unilever demonstrate the effectiveness of green marketing in driving consumer engagement, enhancing brand loyalty, and achieving positive environmental impact. By adopting best practices such as lifecycle assessments, circular economy principles, and employee engagement, businesses can further strengthen their sustainability efforts and position themselves as leaders in their industries. As consumer awareness of environmental issues continues to grow, the importance of green and sustainable marketing will only increase. By prioritizing sustainability, businesses can not only drive business growth and profitability but also contribute to a more sustainable and equitable future for generations to come. Through collective action and ongoing innovation, green marketing can play a pivotal role in shaping a more sustainable economy and society.

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 and progress. These reports provide in-depth information on their circular economy practices and recycling
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RESEARCH ARTICLE

Effect of Alccofine on Water Absorption and Sorptivity of Engineered Cementitious Composites Containing Mono and Hybrid Synthetic **Fibres**

R.Prashanthi^{1*}, K.Sakthimurugan² and G.Pugazhmani³ and S.Gayathri⁴

¹Research Scholar, Department of Civil and Structural Engineering, Annamalai University, Annamalai Nagar, Tamil Nadu, India.

²Professor, Department of Civil Engineering, CK College of Engineering and Technology, Cuddalore, Tamil Nadu, India.

³Assistant Professor, Department of Civil Engineering, Mohamed Sathak Engineering College,) Kilakarai, Tamil Nadu, India.

⁴Assistant Professor, Department of Civil Engineering, Bharath Institute of Engineering and Technology, Hyderabad, India.

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*Address for Correspondence

R.Prashanthi

Research Scholar,

Department of Civil and Structural Engineering,

Annamalai University,

Annamalai Nagar, Tamil Nadu, India.

Email: prashu1691@gmail.com



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ABSTRACT

Water ingression has major influence on durability of concrete materials. This paper deals with a study on influence of Alccofine and synthetic fibers on sorptivity and water absorption in ECC and Hybrid ECC. The sorptivity test measures the rate of movement of water through the concrete under capillary suction as per ASTM C 1585. Water absorption, density and volume of permeable pores in concrete were measured based on ASTM C 642. Based on the literatures, mix design was adopted and fly ash has been replaced by Alccofine and for the workability viscosity modifying agent was used at 0.6% of the binder for all the mixtures. Two types of synthetic fibers, uncoated polyvinyl alcohol (PVA) and polypropylene (PP) were used. The results show that, both ECC and HECC with Alccofine show lower absorption and not much variation in sorptivity than conventional concrete.

Keywords: Engineered Cementitious Composites, Polyvinyl alcohol, Polypropylene, Absorption, Sorptivity





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INTRODUCTION

Concrete should withstand without any deterioration over a design period of the structure. Such concrete is known as durable concrete. The life of concrete may deteriorate when foreign components ingress into it especially water. Durability of the composites mainly depends on the ingression of fluid particles into it, which contains chemical components have ability to deteriorate the composite materials when react with the interfacial transition zone compounds. Lowering the permeability of composite members will eventually reduce penetration of water, chloride ions, sulphate ions, and other harmful substances [1]. After the hydration reaction of cement results a product containing solid with few pore system. The networks of pore system consist of cement paste and allow the fluid to transport into the solid concrete member. The transportation of fluid depends on different factors such as ingredients and proportions of the composite[2]. Sorptivity is an index to measure the moisture transport in the unsaturated specimens [3]. In sorpitivity test water ingression occurs due to the capillary suction in the pore spaces of the concrete not by the pressure head [4]. Sorptivity coefficient is an important index to find the service life of the structure and also to enhance the performance of the composite [5]. Ingression of moisture through capillary suction leads to the carbonation reaction and eventually it deteriorate the composite structure. Replacing the cement by nano materials eventually reduced water absorption and sorptivity in the concrete materials [6]. ECC is a strain hardening cementitious composite designed based on micromechanics and showing high tensile strain capacity and tight crack width control. ECC also possess self healing ability which reduces the deterioration and become a durable material [7]. Quality of concrete is not only based on strength and also based on its durability parameters [8]. Water absorption and sorptivity tests are the easiest method to find the ability of the material to absorb and transmit water by capillarity [9]. Use of glass powder as an aggregate in ECC mixtures increases the permeability properties [10]. Using fibers, mineral admixtures such as flyash, GGBS, silica fume etc improves the performance of the concrete such as toughness, strength as well as durability criteria's [11]. The sorptivity coefficient of self healed ECC also shows better results compared to the preloaded ECC specimens and it shows poorer results when undergo more strain [12]. Water absorption and sorptivity of the concrete with hypo sludge as a cement replacement shows higher values than normal concrete [13]. An acceptance criterion for the durability index has been shown in below table 1 [14].

Water absorption and sorptivity rate of ECC shows better results when compared to the normal concrete [15]. Water absorption in ECC can be controlled by addition of water repellent agent [16]. ECC with rice husk ash shows a higher absorption and void content compared to the ECC with fly ash [17]. Since the crack width is limited to $100~\mu m$, no external ingression of foreign materials is not possible which eventually increases the life of the ECC member [18]. Compared to basalt fiber reinforced concrete, glass fiber reinforced concrete shows higher sorptivity and percentage of voids, both had been tested based on ASTM C 642 and ASTM C 1585 [19]. Sorptivity test gives important information about the pore structure of the composites, tortuosity, continuous capillaries and pore sizes [20]. Addition of fiber in concrete reduce sorptivity in concrete and the reduction is upto 25.85% than control mix [21]. Hence an attempt is made to study the effect of Alccofine and synthetic fibers on the engineered cementitious composites and Hybrid Engineered cementitious composites.

MATERIALS AND METHODS

Alccofine

Alccofine 1203 (Figure 1) is a proprietary patented product (IP Patent No. 297735) with low calcium silicate based mineral additive. Controlled granulation process results in unique particle size distribution. Its latent hydraulic property and pozzolanic reactivity results in enhanced hydration process. Addition of Alccofine 1203 improves the packing density of paste component. This results in lowering water demand, admixture dosage and hence improving strength and durability parameters of concrete at all ages. Fine balance of CaO₂ SiO₂ and Al₂O₃ combined with





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unique patented PSD (Particle Size Distribution) design makes Alccofine a favorable SCM combination for use in all grade concrete.

Cement

Cement used in this work was OPC 53 grade conforming to IS: 8112 – 1989. The chemical properties of cement are given in Table 2. Physical properties of the cement are given in Table 3.

Sand

The fine aggregate available naturally from river beds is used. As per IS 383-1970 Code conforming that, it is coming under Zone III. The particle size distribution curve for fine aggregate is shown in Figure 2. The Specific gravity of sand was found using specific gravity bottle as 2.61

Synthetic Fibers

Synthetic fibres are made only from polymers found in natural gas and the by-products of petroleum. Synthetic fibres are man-made fibres, most of them are prepared from raw material called petrochemicals. All fabrics are obtained from fibres, while fibres are obtained from artificial or man-made sources. Out of these fibers Polyvinyl alcohol fiber and polypropylene fibers were used in this study.

Polyvinyl Alcohol (PVA) Fiber

Polyvinyl Alcohol is the main raw material to produce PVA fiber. It undergoes process of dissolution, spinning, heat – setting, cutting, and balling to form a high strength, high modulus fiber. The physical properties of PVA fiber was shown in Table 4. Figure 3 shows the PVA fiber which was used in this study.

Polypropylene (PP) Fiber

Polypropylene fiber is a kind of linear polymer synthetic fiber obtained from propylene polymerization. It has some advantages such as light weight, high strength, high toughness and corrosion resistance. The physical properties of PP fiber was shown in Table 5. Figure 3 shows the PP fiber which was used in this study.

Sika Viscocrete

Sika viscocrete is a polycarboxylate based high performance super plasticizing admixture which imparts high workability, prolonged workability retention and allows a large reduction in water content. It produces a more uniformly cohesive high quality free flowing concrete. Specific gravity of viscocrete is 1.08. Figure 4 shows the admixture can.

Mix Proportions

Snce it is an evolving material no proper mix designs were available. Based on the literatures as a reference, mix design was taken as per trial and error method. Totally 6 mixes were prepared. They are control mix CC, ECC-2%PVA+0%PP (M₁), ECC-0%PVA+2%PP (M₂), HECC-1%PVA+1%PP (M₃), HECC-1.5%PVA+0.5%PP (M₄), HECC-0.5%PVA+1.5%PP (M₅). Mix design used in this study is given in Table 6 along with its respective compressive strength. Simplified mix ratio is given in Table 7.

Mixing and Specimen Preparation

Heavy duty hand driller with mixing bit was used to cast both control and ECC specimens. For each mix 3 numbers of 70.7mm cubes for water absorption and 3 numbers of 100mm diameter 50mm height sorptivity specimens were cast. For both control and ECC, same method of mixing was taken place to ensure uniform dispersion of fibers in the matrix since it plays a major role in the performance. Cement, Alccofine and sand were dry mixed till it gets uniform colour. After few minutes of mixing, water along with sika viscocrete was added and mixed with high speed for





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about 5 minutes. Once the uniformity attained, fibers were added randomly and mixed for 3 more minutes or till the uniformity achieved. After the completion of mixing the slurry was made to pour into the oiled moulds and kept for 24 hrs in the normal temperature. Specimens were removed from the moulds after 24 hrs and kept in the water curing for 28 days. The number of specimens casted was shown in the Table 8.

Experimental Works

Absorption after Immersion

Water absorption after immersion is the amount of water absorbed by the specimen after immersing the oven dried specimen in the water for about 48 hrs. The test was performed based on ASTM C 642. Figure 5 shows the cubes casted and cured, specimens dried in the oven, specimens immersed in water.

Absorption after Immersion and boiling

Absorption after immersion and boiling can be found by immersing the specimens in the boiling water for 5hrs and allow it to cool for 14hrs naturally and weight has to be taken by removing the surface moisture by wiping the specimen by dry towel. Figure 6(a) shows the specimens in the boiling water and 6(b) shows measuring apparent weight of the specimen.

Bulk Density (Dry)

Dry bulk density is defined as the mass of the dry specimen divided by the total volume of the wet sample. It is given by below formula,

Bulk Density = $\frac{oven\ dry\ mass}{(Saturated\ mass\ after\ boiling\ -immersed\ apparent\ mass\)}*$ density of water

Bulk Density after Immersion

Bulk density after immersion is the ratio of mass of the immersed specimen by the total volume of the wet sample. It is given by below formula,

Bulk density after immersion = $\frac{Saturated \ mass \ after \ immersion}{(sat \ mass \ after \ boiling - immersed \ app \ mass)}^* density of water$

Bulk Density after Immersion and Boiling

Bulk density after immersion and boiling is the ratio of mass of saturated mass after boiling by the total volume of the wet sample. It is given by below formula,

the wet sample. It is given by below formula, $Density after immersion and boiling = \frac{Saturated mass after boiling}{(Sat mass after boiling - immersed app mass)} * density of water$

Apparent Density

Apparent density is the ratio of oven dry mass to the difference between oven dry mass and immersed apparent mass. It is given by below formula,

Apparent Density = $\frac{oven \ dry \ mass}{(oven \ dry \ mass - Immersed \ apparent \ mass)} * density of water$

Volume of permeable pore space (Voids)

Out of entire cement paste one third to one half become interface transition zone. Since it is the locus point for cracking highly prone to become porous but permeability of the concrete is depends on the bulk portion of the hardened cement paste since it is the continuous phase (A.M.Neville, 2011). Volume of permeable pore space (voids) can be found by the ratio of difference of bulk and apparent density to apparent density. It is given by below formula,

Voids (%) = $\frac{(apparent\ density\ -bulk\ density\)}{apparent\ density} * 100$

Sorptivity

Sorption in the concrete member is due to the capillary suction through the pores. Since the capillary suction would not takes place in either dry or saturated concrete. Sorptivity test specimens were oven dried for at least 48hrs after 28 days of water curing to remove all the moisture from the concrete specimens. After removing from the oven it has to covered tightly inside polythene bags for about 15 days. After removing the specimens from the bags, leaving one





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side exposed to water all other sides were covered with the electrician insulation tape and placed in the sorptivity setup up keeping 3 – 5 mm depth of water as per ASTM C 1585. This test method is used to find the rate of absorption of water by the cementitious specimens by measuring the mass gain in the specimen resulting from the sorption of water as a function of time when only one surface exposed to the water. This test was performed based on ASTM C 1585. Figure 7(a) shows diagrammatic representation of sorptivity setup, 7(b) shows the specimens prepared for sorptivity test, 7(c) shows sorptivity test setup, 7(d) shows sorptivity setup with water. The absorption was calculated by below given formulas,

RESULTS AND DISCUSSIONS

Absorption after Immersion

The test results were shown in the bar chart in figure 8. From the figure it is observed that all the mixes showed lower water absorption that control mix, it means that Alccofine plays a major role in the pore refinement. When using a single type of fiber in ECC does not create any impact on water absorption but in the hybrid ECC, mix with 1.5% PVA and 0.5% PP shows a relatively lesser absorption than mixes containing 1%, 0.5% PVA and 1%, 1.5% PP. Since PVA is a high modulus fiber and occurrence of agglomeration is comparatively lower than PP fibers. While using 1.5% PVA and 0.5% PP occurrence of agglomeration is relatively low and leads to the lesser water absorption. Absorption of water is relatively high when using poly propylene fiber compared to the steel fibers, glass fibers, waste plastic fibers and high density poly ethylene fibers (M. P, 2019) and it reflects in the hybridization of polypropylene fiber with polyvinyl alcohol fiber. However poly vinyl alcohol fiber showed lesser absorption. It can be found that 28.52% absorption in M_1 , 12.93% absorption in M_2 , 17.87% absorption in M_3 , 28.14% absorption in M_4 , 21.29% absorption in M_5 had been reduced compared to the control mix.

Absorption after Immersion and boiling

The test results were shown in the bar chart in figure 9. From the figure it is observed that all the mixes showed lower water absorption than control mix even after boiling, it means that Alccofine plays a major role in the pore refinement. Synthetic fibers are prone to melt when the temperature rises but in the boiling stage outer surface of composites act as a non porous material because of pore refinement occurred and boiling water does not affect the synthetic fibers inside the specimen. Hence it shows similar trend of absorption in both normal and boiling water. However all the ECC mixes shows lesser water absorption than control mortar.

Bulk Density (Dry)

The test results were shown in the figure 10. Compared to the control mortar both mono and hybrid ECC shows higher density and that is because of the addition of Alccofine. But there is a slight variations in the density of the hybrid ECC compared to the Mono fiber ECC and that is may be because of the differences in the fiber texture and their compatibility. Though there is differences in the density since it is in less amount it can be neglected. It can be found that density of 22.83% in M_1 , 24.41% in M_2 , 11.81% in M_3 , 19.69% in M_4 , 18.11% in M_5 had been increased compared to the control mix.





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Bulk Density after Immersion

The test results were shown in the figure 11. Compared to dry density, density after immersion shows no significant differences whereas based on the absorption, density had changed. Density increases with increase in volumetric moisture content (zha, 2018). As per literature there will be rise in density whenever the moisture content rises. Below figure also showing the same.

Bulk Density after Immersion and Boiling

Since the boiling temperature didn't affect the synthetic fibers there is no significant difference encountered in the density values. After immersion and boiling the density of the specimens remain same as after immersion. Even though the fibers are prone to melt during high temperatures, matrix phase has covered the fibers during boiling. Due to that there is no damage to the fibers and internal structure of the composite.

Apparent Density and Voids

Apparent density is mass per unit volume of the particles, excluding the voids in the material. Relationship between the apparent density and voids is shown in the fig 12. When the voids increased apparent density got reduced. From the graph it is evident that control mix i.e without Alccofine and fibers shows higher percentage of voids whereas mixes having Alccofine and fibers showing relatively lesser voids. Based on the percentage of the voids apparent density also get altered. Control mix showing lesser density compared to other mixes since the percentage of voids are more in it compared to the other mixes which containing Alccofine and fibers. Both ECC and HECC shows almost same percentage of voids and it resulted in the apparent density also. It can be found that 27.44% voids in M₁, 28.28% voids in M₂, 24.92% voids in M₃, 25.25% voids in M₄, 23.57% voids in M₅ had been reduced compared to the control mix.

Sorptivity

After the sorptivity test it was observed that both ECC and HECC shows relatively higer capillary suction in the intial time period that is upto six hours and eventually got reduced when it entered the secondary absorption. Whereas the control mix shows relatively lesser absorption in the intial period of absorption and it is more compared to ECC and HECC in the secondary stage of absorption. This happens because of the change in the continous pores in the ECC and HECC. Because of the presence of fibre in high percentage both ECC and HECC allows the suction in intial time period but it has got eventually reduce when it comes to the secondary sorption. This is because fibres obstruct the path randomly and it changes the totuosity of the specimen. However control mix shows lesser absorption in the intial period of time increased slowly than ECC and HECC in the secondary capillary suction. It happened due to the absence of the fibre and alccofine whereas Alccofine plays a major role in the pore refinement and fiber become the responsibility for the change in the tortuosity of the specimen. Upto 2% addition of fibers did not affect the sorptivity property of the specimens whereas when it increase to 3% the sorptivity value in both inital and secondary will increase [22]. Figure 13 shows sorptivity curves of control and ECC. Figure 14 shows sorptivity curves of control and HECC. From the graph it is inference that initial absorption of all the mixes including control mix show more or less similar trend whereas after 6 hrs both ECC and HECC showed lesser capillary suction than the control mix. Eventhough pore refinement occurs because of the Alccofine, presence of fibers altering the tortuosity of the specimens. However PVA fibre shows higher suction than PP fibres it is because of their specific gravity. Since PVA is a high modulus fibre for the given V_f quantity is little lesser than polypropylene. The initial and secondary sorptivity of all mixes are tabulated in table 9.

CONCLUSION

From the above experimental tests carried out the following conclusions have made, Water absorption of ECC and HECC specimens showed better results compared to the control specimen. Whereas specimens with polypropylene fibers show relatively higher water absorption compared to the specimens containing poly vinyl alcohol fiber. Since the temperature is not high while boiling the water for 5 hours when the specimens immersed in it, didn't cause any





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damages to the fibers present inside the specimens. The absorption capacity did not vary much after boiling the specimens. Regarding bulk density of control, ECC and HECC, ECC shows higher density than control and HECC. The slight reduction of density in the HECC is lack of compatibility of the fibers but the difference is very less than ECC compared to the control specimen. Bulk density after immersion shows variations in its value since it is depends on the moisture content. As per the test it shows that the bulk density after immersion is proportional to its moisture content and both ECC and HECC show favourable results. Since Alcofine plays a major role in the pore refinement compared to the control specimens ECC and HECC shows lesser percentage of voids. Because of the lesser voids in the ECC and HECC apparent density is higher compared to the control mix. Sorptivity values of the control specimen shows lesser absorption than ECC and HECC in the initial period that is upto six hours whereas after that ECC and HECC shows lower soprtivity than control in the secondary phase. It is because of fibers which change the tortuosity of the specimens which in turn reduces the sorptivity in ECC and HECC.

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Table 1. Acceptance Criteria for Durability Indexes

Acceptance Criteria	Oxygen Permeability Index (OPI) (log scale)	Sorptivity (mm/h)
Workroom concrete	> 10	< 6
As-built Structures		
100% recognition	> 9.4	< 9
50% recognition	9.0 to 9.4	9 to 12
Helpful measures	8.75 to 9.0	12 to 15
Elimination	< 8. 75	> 15

Table 2. The chemical Analysis of Cement

Chemical composition, %	SiO ₂	CaO	Al ₂ O ₃	Fe ₂ O ₃	MgO	K ₂ O	SO ₃	Na ₂ O	LOI
Cement	21.80	63.56	5.12	3.20	0.80	0.75	3.22	0.55	1.00

Table 3. The Physical Properties of Cement

Sl No	Properties	Values
1.	Specific Gravity	3.10
2.	Specific Surface Area	3710 cm ² /g
3.	Normal Consistency	31%
4.	Initial Setting Time	60 mins
5.	Final Setting Time	330 mins

Table 4. The Physical Properties of PVA fiber

Туре	Fiber Diameter (µm)	Length (mm)	Specific Gravity	Tensile Strength (MPa)	Elongation (%)	Young's Modulus (GPa)
PVA	30	12	1.3	1700	6	40

Table 5. The Physical Properties of PP fiber

Type	Fiber Diameter (µm)	Length (mm)	Specific Gravity	Tensile Strength (MPa)	Elongation (%)	Young's Modulus (GPa)
PP	30	12	0.91	550	25	5

Table 6. Mix Proportions

Mix	Comont	Alccofine	Fine	Water	Fibers ((kg/m³)	HRWR	Compressive
ID	Cement (kg/m³)	(kg/m³)	Aggregate (kg/m³)	(kg/m³)	PVA	PP	(kg/m³)	Strength (N/mm²)





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CC	640.2	514.8	574	310	-	-	7.02	58.06
M ₁	640.2	514.8	574	310	24	-	7.02	76.44
M ₂	640.2	514.8	574	310	-	24	7.02	74.14
M 3	640.2	514.8	574	310	12	12	7.02	71.58
M ₄	640.2	514.8	574	310	18	6	7.02	72.44
M ₅	640.2	514.8	574	310	6	18	7.02	75.32

Table 7. Mix Ratio

Cement	Alccofine	S/B	W/B	Fiber	HRWR
1	0.8	0.5 0.27	2% of	0.6% of	
1			0.27	binder	binder

Table 8. Number of Specimens

Sl No	Specimen	No. of Specimens	Type of Test	
1	Cube (70.7mm)	6 x 3 = 18	Water absorption as per	ASTM C 642
2	Cylinder (100mm dia, 50 mm height)	6 x 3 =18	Sorptivity as per	ASTM C 1585

Table 9. Initial and secondary sorptivity of all mixes

Tuble 3. Illitial and secondary surprivity of all hirkes					
Sl No	Mix ID	Average initial sorptivity in 10 ⁻³ x	Average secondary sorptivity in 10-		
		mm/sec ^{1/2}	³ x mm/sec ^{1/2}		
1	M_1	20.4	5.15		
2	M ₂	20.14	5.15		
3	M 3	20.80	4.99		
4	M_4	21.47	4.35		
5	M ₅	21.67	4.99		
6	Control	24.99	5.93		



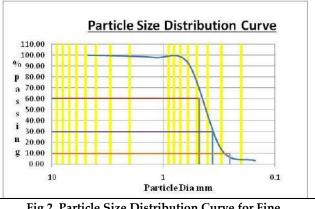


Fig.1 Alccofine

Fig 2. Particle Size Distribution Curve for Fine Aggregate





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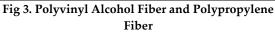




Fig 4. Sika Visco Crete









Figure 5 Cubes Casted and in Curing, Specimens dried in oven, Specimens immersed in water





Figure 6(a) Specimens in the boiling water (b) Measuring apparent weight of the specimen

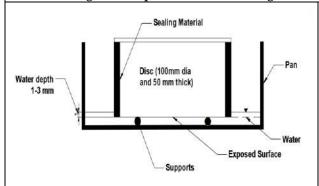


Figure 7(a). Diagrammatic representation of sorptivity setup



Figure 7(b). Specimens prepared for sorptivity test





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Figure 7(c) Sorptivity test setup

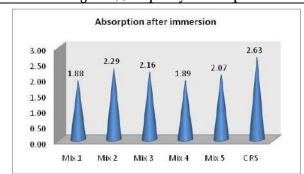


Figure 7(d) Test setup with water

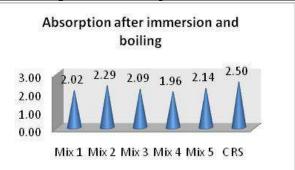


Figure 8. Water absorption after immersion

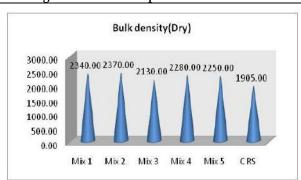


Figure 9. Water absorption after immersion and boiling

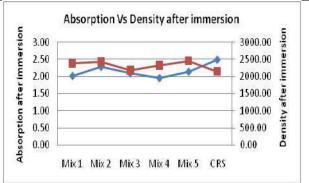


Figure 10. Dry bulk density

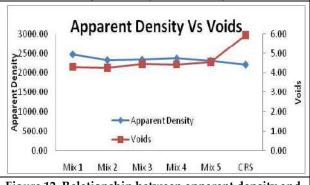


Figure 11. Water absorption Vs Density after immersion

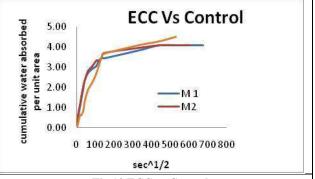
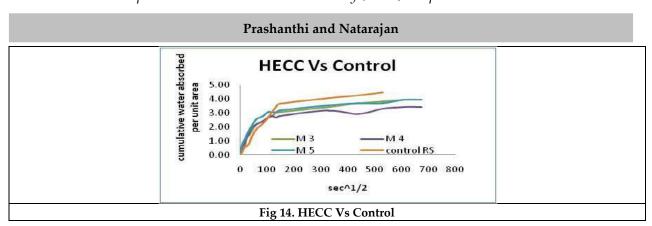


Figure 12. Relationship between apparent density and voids

Fig 13 ECC vs Control











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RESEARCH ARTICLE

A Study on the Urine Analysis of Selected Teenagers using Dipstick Method

T. Amala Arockia Raj¹, M.Muniya Naik², K.Janardhana³, A.Govardhan Naik⁴, A. Jayasankar⁵, D. Veera Nagendra Kumar⁶ and V. UdayKiran^{7*}

¹Principal and Lecturer, Department of Chemistry, Loyola Degree College (YSRR) -Pulivendula (Affiliated to Yogi Vemana University, Kadapa) Andhra Pradesh, India.

²Lecturer, Department of Zoology, Government Degree College, Rayachoty (Affilated to Yogi Vemana University, Kadapa) Andhra Pradesh, India.

³Lecturer, Department of Zoology, Government Degree College, Rajampeta (Affiliated to Yogi Vemana University, Kadapa) Andhra Pradesh, India.

⁴Teaching Assistant, Department of Zoology, SVU College of Sciences, Sri Venkateswara University, (Affiliated to Sri Venkateswara University) Tirupati, Andhra Pradesh, India.

⁵Lecturer, Department of Zoology, S.V.C.R. Government Degree College, Palamaner, Chittoor (Affiliated to Sri Venkateswara University Tirupati) Andhra Pradesh, India.

⁶Lecturer, Department of Zoology, Government College for Men (A), (Affiliated to Yogi Vemana University) Kadapa, Andhra Pradesh, India.

Lecturer, Department of Zoology, Loyola Degree College (YSRR) -Pulivendula (Affiliated to Yogi Vemana University, Kadapa) Andhra Pradesh, India.

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*Address for Correspondence

V. UdayKiran

Lecturer,

Department of Zoology,

Loyola Degree College (YSRR) –Pulivendula (Affiliated to Yogi Vemana University, Kadapa) Andhra Pradesh, India.

Email: vempati.uday6@gmail.com



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ABSTRACT

Urine is frequently used for clinical diagnostics and biomedical research. Urine analysis has been studied and used since ancient times. Urine contains a diverse set of metabolites that can provide information about the body's current physiologic condition as well as clinical signs of disease. We tried to explore the benefits and limits of the urine dipstick as a low-cost tool in point-of-care settings, as well as the reasons for its lack of use as a broad screening tool. Commercially available dipsticks can be used to measure urine pH and various contents (e.g., leukocytes, erythrocytes, protein, glucose, nitrate, ketones, blood, bilirubin, urobilinogen) as a kind of fast method for diagnosis. It is a quick, uncomplicated process indicates any abnormal human health conditions and allows the removal of unwanted elements such as





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those significantly polluted by microbial infection or blood. However, the use of dipsticks for inclusion/exclusion in urine research has been quite debatable to date: there is no common agreement on tests for which dipstick analysis is best or where to set inclusion/exclusion criteria.

Keywords: urinalysis, dipstick, dry chemistry, urine, assay principles.

INTRODUCTION

Urine, often regarded as a liquid goldmine of diagnostic information, has been employed as a diagnostic medium for centuries. Its analysis has evolved from the rudimentary observations of color and odor to sophisticated laboratory techniques that enable the detection of specific biomarkers indicative of various medical conditions(Decavele A-SCet al.,2012). Among the plethora of valuable substances present in urine, albumin and glucose stand out as critical indicators of renal and metabolic health. Monitoring these components through dipstick analysis of urine chemistry offers a rapid and cost-effective means of assessing a patient's condition, making it a vital tool in both clinical and research settings(Delanghe Jet al.,2014). Albuminuria, the presence of excess albumin in urine, is a well-established marker for kidney dysfunction and a harbinger of renal disease progression. It plays a pivotal role in identifying patients at risk for various renal conditions, such as diabetic nephropathy and hypertensive nephrosclerosis (Ercan Met al.,2015). On the other hand, glycosuria, the presence of glucose in urine, is a key diagnostic criterion for diabetes mellitus and provides valuable insights into glycemic control (Haber MH (2010)).

Monitoring these markers through dipstick analysis offers advantages of simplicity, speed, and cost-efficiency, making it an attractive option for routine diagnostic evaluations (Herrington*et al.*, 2016). In this research article, we delve into the diagnostic evaluation of albumin and glucose in urine using dipstick analysis, exploring the principles behind this technique, its clinical applications, and its potential limitations (LaRocco MT, *et al.*,2016). We aim to provide a comprehensive overview of the state-of-the-art in dipstick analysis for albumin and glucose, including advancements in technology and methodologies (Pugia MJ (2000). Additionally, we will discuss the clinical significance of dipstick analysis results, offering insights into how this technique can aid in the early detection and management of renal and metabolic disorders (Rotter M, et al (2017). As the demand for rapid, point-of-care diagnostic tools continues to grow, understanding the utility and reliability of dipstick analysis for albumin and glucose becomes paramount (DelangheJR*et al.*,2017). This research article endeavors to shed light on the current landscape of dipstick-based urine chemistry analysis, empowering healthcare professionals, researchers, and clinicians with the knowledge needed to make informed decisions in the realm of diagnostic medicine (Sureda-Vives M, *et al.*,2017).

METHODS AND MATERIALS

TEST STRIP TECHNOLOGY

The analytical sensitivity of urine test strips has increased due to advancements in electronic detection, allowing for the quantitative examination of red blood cells, white blood cells, glucose, and urinary protein (Oyaert, *et al*,.2018). This has been used to analyze albuminuria quantitatively and determine the albumin: creatinine ratio in the micro albuminuria range (20-200 mg/L) for both ketones and albumin. The albuminuria measurement can be assisted by the creatinine-specific test pad, which corrects urinary dilution. Leukocyte esterase and peroxidase activity readings are sensitive because to CMOS technology. Sensitive dyes have increased sensitivity. Results from urine test strips are now read and interpreted using smart phones (Oyaert MN, *et al* 2018). Urine dipstick dry chemistry is a low-cost, simple approach for evaluating numerous assay principles in a short period of time, making it excellent for use at point-of-care settings. However, storage conditions must be carefully monitored to avoid microbe and metabolite





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overgrowth. The test can measure pH, osmolality, hemoglobin/myoglobin/hematuria, leukocyte esterase, glucose, proteinuria, nitrites, ketones, and bilirubin, which can be used to guide further testing if clinical suspicion exists. Dipsticks can also be automated, reducing human error and labour and making them suited for point-of-care applications. They do, however, have drawbacks, including a lack of measurable data, false positives and negatives due to confounding factors, and interpretation uncertainty. Furthermore, dipsticks do not have enough sensitivity for albumin in normal home testing settings, and alternative markers, such as the albumin-to-creatinine ratio, are more useful for diagnosing diabetes-associated microalbuminuria.

Sample Collection and Preparation

Of the 50 candidates identified for the this test, we had chosen only 18 samples after discriminating based on the Body weight and past medical history. The samples for this test were having body weight below 50Kgs and some health issues. They were of the age group of between 19-20 years. A fresh urine sample was collected in a clean and dry container. It was not centrifuged. The sample was well mixed before taking the test. The urine test was conducted within an hour. All specimens were always taken and kept under sanitary conditions.

Points for Attention*Proteinandmicroalbumin* (10VOnly)

Protein and microalbumin are more sensitive to albumin than Bence-Jones proteins, immune globulin and haemoglobin. When it is negative, it doesn't mean that protein does not exist.

Glucose

The test is only for glucose. Other than glucose, no other component in urine has been shown to produce a positive result. 2.2mmol/L glucose in 0.28mmol/L ascorbic acid-containing urine may cause a colour shift that could be perceived as positive. Concentrations of ascorbic acid as low as 0.28mmol/L and/or acetoacetic acid as low as 1.1mmol/L may not affect the test. A tiny amount of glucose is normally eliminated by the kidney. Typically, the amount is less than the sensitivity of the reagent test.

Bilirubin

Bilirubin in urine is typically impossible to detect using even the most sensitive approach. A closer look is necessary if there is little bilirubin present in the urine. Any substance that turns an acidic medium red, such as phenazopyridine, or medications that dye urine red may have an impact on the test's outcome. A false negative result could occur from a high ascorbic acid content.

Ketone

Acetoacetic acid in urine causes the reagent strip to respond. Neither acetone nor ß-hydrobutyric acid are compatible with it. Normal urine samples typically yield negative test findings. Highly pigmented urine or urine that contains a significant amount of levodopa metabolites may yield false-positive results.

Specific Gravity

Urine samples can have a specific gravity between 1.000 and 1.030 with a mean error of 0.005 when using the reagent strip for specific gravity. pH values, non-ionic components, highly buffered alkaline urine, and moderate protein concentrations can all be altered. Instruments used to measure urine automatically correct strip readings(https://www.nu-careproducts.co.uk/data.).

Blood

Patients may react differently to the "trace." Individual cases necessitate clinical judgment. The need for additional diagnostic testing is indicated by the appearance of green specks (intact erythrocytes) or green colour (haemoglobin/myglobin) on the reagent area within 60 seconds. The urine of women who are menstruation frequently contains blood. About 5–15 undamaged erythrocytes per litre of haemoglobin, which ranges from 150–620 g/L. The reagent strip can be used in addition to the microscopic inspection because it is particularly sensitive to haemoglobin. In urine with a high specific gravity, the strip's sensitivity may be diminished. Both myglobin





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and haemoglobin are equally sensitive to the strips. A few oxidising impurities, such hypochlorite, can produce erroneous positive findings. It's also possible for microbial peroxidase linked to urinary tract infections to result in a false-positive test. Less than 5.Ommol/L of ascorbic acid in the urine may not affect the test's outcome (**Wilson MLet al., 2004**).

pΗ

The pH test are a measures pH values in the range of 5.0-9.0 instrumentally and 5.0-8.5 visually.

Urobilinogen

Urobilinogen can be found in urine in amounts as low as 3 mol/L (or roughly 0.02 Ehrlich unit/dL) using the reagent strips. Urine results of 33 mol/L show the crucial value, signifying the change from normal to abnormal, necessitating additional examination of the patients and specimens. The absence of urobilinogen cannot be definitively determined by the negative results.

Nitrite

Nitrite, which comes from food-derived nitrate, is produced in the urine by gram-negative bacteria. Since other elements in urine will not react with the reagent strip, nitrite is the only element that can be detected. Positive results should not be extrapolated from pink patches or edges on the strip. However, the development of any consistent pink colour in any degree should be seen favourably. There is no direct correlation between the levels of colour development and bacterial populations. The absence of germs in high quantities is not implied by the bad outcome (**Penders J, et al, 2002**).

Negative results may occur

• The nitrate in the foods is absent. Large high volume of specific gravity in urine may reduce the sensitivity of the test. 2.8mmol/Lascorbicacid or lesswon'tin therefore with the test result.

Leukocytes

Leucocytes (granulocytic leukocytes) tested respond with esterase. Normal urine samples often produce negative findings; clinically significant positive values are (+ or more). 'Trace' outcomes that are detected once or twice might not have much clinical importance, whereas results that are observed regularly might. Random samples from females may occasionally yield "positive" findings because vaginal discharge contaminated the material. Test findings may be affected by high specific gravity or elevated glucose levels (over 160 mmol/L).

Protein

Since the reagent region is sensitive to albumin, a negative result in urine tests is insufficient to prove the presence of proteins such as globulins, haemoglobin, Bence Jones protein, and mucoprotein. False positive results can occur in highly buffered alkaline urines (Lifshitz E, Kramer L (2000)).

Interpretation of Results

We collected 50 samples for the study, but only 18 of them showed positive and pertinent results. Since the majority of the samples are from teenagers (0.0 mmoI/L), their metabolic rate is significantly correct, we discover that all values are strongly negative. Out of the 18 nitrate samples that were discovered, 13 showed positive results, with the highest sample reaching 70 nmoI/L. Ketone levels are significantly abnormal when compared to other parameters, with 6 samples out of 18 being abnormal (it turns purple color: highest levels 16.0 mmoI/L), two samples being abnormal (3.9 mmoI/L), five samples being abnormal (0.5 mmoI/L), one sample being abnormal (1.5 mmoI/L), and three samples being negative. In the meanwhile, the blood parameters were analysed with urine sample, which was significantly traced with 15 samples out of 18 identified ones.Out of 18 samples, only one has a bilirubin parameter that is negative, while the other 17 are all significantly positive, with the highest value being 51 µmoI/L it turns in orange color and the lowest value being 17µmoI/L.Two samples had the highest amounts of positive protein (0.3





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g/L), and five protein readings are normal. 11 other samples are noticeably positive. Only three of the 18 samples used in this study's estimation of uribilonogen levels revealed negative results, which meant that the pH, specific gravity values and the numbers of leucocytes were most likely aberrant.

Overview of urine chemical composition analysis using dipsticks and assay concepts

The process of detecting urine on a dipstick and comparing the colour change to a reference standard is known as dipstick/dry chemistry analysis (DelangheJ R et al, 2016). A chromogenic chemical often reacts with another product to cause this. Binding dye compounds, enzymatic reactions, immunologic reactions, and catalysis of oxidationreduction reactions are examples of common reactions (Dolscheid - Pommerich RC et al, 2016). The majority of analyses rely on colorimetry, with the intensity of colour change proportional to analyte concentration. Due to their comfort, accessibility, and accuracy, dipstick readers have become a common tool in primary care, hospitals, and the consumer market (Echeverry G, et al, 2010). Despite the possibility of imprecise evaluations based on visual acuity and colour vision, these devices convert photonic energy from coloured reaction products into electrical signals, allowing the generation of semi-quantitative/quantitative data. Dipstick/dry chemistry is used in automated systems such as the Beckman Coulter IRICELLTM to maximize specimen throughput, minimize handling, and reduce manual labour (Fogazzi GB et al, 2008). These technologies have been well researched for application in dipstick interpretation. Dipstick readers with a 'lab on a chip' are also available for smart phone use. Dipstick/dry chemistry analysis can provide useful information regarding the qualities of urine samples, such as osmolarity (Frazee BW et al, 2012). Sample pH can be calculated within a range of 5.0-8.5, with variations possible at the extremes (5.5 and >7.5). The use of a glass electrode is the preferred and most accurate approach for measuring pH at those values (Hall JE (2016).

CONCLUSION

This is a preliminary study to expose the students towards analytical tools and to make them aware of the health parameters. Those who were with high protein, Ketone and Nitrate levels were directed for further medical checkup. Those who had blood cell damage were instructed to for further treatment to improve their blood quality.

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Authors' Disclosures of Potential Conflicts of Interest

No potential conflicts of interest relevant to this article are reported.

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Table Detection of several parameters in urine using the dipstick method through urine analysis

			Name of parameters with respected values Using H10 Reagent Strips for Urinalysis (Dip-stick method)										
S.no	C = a C = = 1 a		Specific	Glucose	Glucose Leucocytes		Protein	Blood	Ketone	Bilirubin	Urobilonogen		
5.no Sa	Sample	pН	gravity	(mmoI/L)	(leuko/µL)	(µmoI/L)	(g/L)	(Ery/µL)	(mmoI/L)	(µmoI/L)	(µmoI/L)		
1	S_1	5.0	1.030	0.0 (-)	70	Positive	Positive	Traced	3.9	17	17		
2	S ₂	6.0	1.000	0.0 (-)	00(-)	Negative	Normal	Non- Trac	Negetive	17	Negative		
3	S_3	6.0	1.030	0.0 (-)	70	Positive	Positive	Traced	16	17	17		
4	S_4	6.0	1.030	0.0 (-)	70	Positive	Positive	Traced	16	17	17		
5	S_5	6.0	1.030	0.0 (-)	70	Positive	Positive	Traced	16	17	17		
6	S_6	6.0	1.005	0.0 (-)	70	Positive	Positive	Traced	0.5	17	17		
7	S_7	5.0	1.030	0.0 (-)	15	Positive	Positive	Traced	16	17	3.4		
8	S_8	6.5	1.010	0.0 (-)	0.0 (-)	Negative	Negetive	Non- Trac	0.5	Negative	Negative		
9	S 9	6.0	1.030	0.0 (-)	70	Negative	Positive	Traced	16	17	17		
10	S ₁₀	6.0	1.030	0.0 (-)	70	Positive	Positive	Traced	3.9	17	17		





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11	S ₁₁	5.0	5.0	0.0 (-)	15	Positive	Positive	Traced	1.5	17	17
12	S ₁₂	6.0	1.030	0.0 (-)	70	70 Negative		Traced	0.5	51	17
13	S ₁₃	5.0	7.0	0.0 (-)	70	70	Negetive	Traced	Negative	51	17
14	S ₁₄	6.0	1.030	0.0 (-)	70	Positive	0.3	Traced	16	51	17
15	S ₁₅	6.0	1.025	0.0 (-)	0.15	Positive	Positive	Traced	16	51	17
16	S ₁₆	5.0	1.030	0.0 (-)	70	Positive	Positive	Traced	0.5	17	17
17	S ₁₇	5.0	1.000	0.0 (-)	0.0 (-)	Negative	Normal	Non- Trac	Negative	17	Negative
18	S ₁₈	6.0	1.030	0.0 (-)	70	Positive	Negetive	Traced	0.5	17	17

Green color indicates Negative (-ve), Brown indicates Positive (+ve) and Red indicates severity.

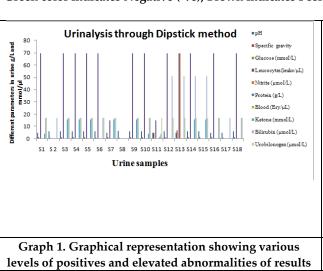






Fig 1: Collection of Samples



Fig 2: Results strip

Fig 3: Strips revealing positive results (+Ve)





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RESEARCH ARTICLE

Automatic Power Factor Correction Unit using Arduino for Smart Homes

Krishnaveni S1*, Marutham Rathna Valli M2, Keerthana S2 and Kaviyamalar A.D2

¹Associate Professor, Department of EEE, Sri Sivasubramaniya Nadar College of Engineering, (Affiliated to Anna University), Chennai, Tamil Nadu, India.

²UG Student, Department of EEE, Sri Sivasubramaniya Nadar College of Engineering, (Affiliated to Anna University), Chennai, Tamil Nadu, India.

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*Address for Correspondence

Krishnaveni S

Associate Professor, Department of EEE, Sri Sivasubramaniya Nadar College of Engineering, (Affiliated to Anna University), Chennai, Tamil Nadu, India. Email: krishnavenis@ssn.edu.in



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ABSTRACT

In order to expedite our everyday routine, the consumption of electrical equipment has greatly expanded in the modern world. However, the majority of customers also depend on their projected monthly bills from the public sector. Therefore, managing power consumption and the associated payment amount requires information of power usage, load type, and appliance management. When inductive loads are used more frequently, the power factor drops to an abnormally low level, drawing more current and increasing the amount of electricity used. Thus, this research discusses a less expensive way to modify the power factor at the home level. This project outlines the process for using capacitor bank sensors to help modify the lagging power factor and Arduino UNO, which is compatible with your home's current setup. This technique improves efficiency, reduces the power factor penalty in small-scale companies, and upgrades and modernizes dwellings with modern technology.

Keywords: Power factor, Real power, Apparent power, Arduino UNO, Capacitor banks

INTRODUCTION

In our era of rapid advancement, energy efficiency is now regarded as one of the main goals. The present technological revolution is trying to fulfill everything needed by the society in a faster pace and so electrical power requirement is so enormous and very precious. TamilNadu Electricity Board [TNEB] tariff is increased every year due to bimonthly payment system since the electricity tariff depending on the conventional power generation. Moreover, the conventional power resources are in decline phase to meet today's population demand and creates the





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alert everywhere how the electrical energy usage has to be curtailed. So, power consumption is superabundant which has a direct impact on the electricity bill, which remains an economic burden in lower and middle-income families. Therefore, the knowledge on electrical energy saving methods must be spread over and the awareness should be created to reduce the electrical usage by the consumer. Many ways are there to minimize the power consumption which includes manual methods where the user has to handle the electrical energy preciously, proper cut off of electrical appliances when not in use, knowing everyday power consumption by SMS alert [1-3], appliances management based on peak hours and maximum demand [4], and installing the power factor correction device. The ultimate idea of introducing various concepts regarding conscious use of electricity and proper maintenance and monitoring of power is to reduce the overall billing cost of the consumer, which has a major social and economic impact in daily lives.

The power quality has also become a hot topic recently due to rapid increase in number of the electronic appliances at home and industries, and the researches in power quality reveals the importance of it. Efficiency is an important measurement on power quality and increasing energy efficiency using power factor correction device has many advantages. Most of the industries are dependent on inductive loads, which on the other hand decreases the power factor and which obviously decreases the efficacy of the system [5]. An Automatic Power factor correction device has been proposed where an Arduino plays a major role at lesser cost is proposed in this paper. This method estimates the phase angle $lag(\emptyset)$ of current and leads to the calculation of the power factor (cos \emptyset). The Arduino calculates the required adjustments to normalize the power factor value. This technique can be implemented in households, small scale industries and in even bigger firms so that the system might become more efficient and stable.

Importance of power factor improvement

The complex power, sometimes called the total power, is the total of the real and reactive powers added together. The entire power is calculated in VA and taken into account when charging for energy. [6].

$$Power factor(pf) = \frac{True Power (W)}{Apparent Power (VA)}$$
(1)

Figure 1 shows the power triangle where the angle between the real power and the total power is defined as the power factor angle φ . The electrical load may have a power factor of less than one if reactive power results in the real power being less than the perceived power. Reactive power deliberately raises current in power companies, which results in worse operational efficacy and lost income. Therefore, power sectors mandate that their customers maintain their power factors over a certain threshold in order to avoid additional penalty. Reactive power may be due to connecting wires, circuit breakers, switches, power transformers, and transmission lines with higher current capacity. So, the lower power factor in turns into lower efficiency, large copper loss, greater conductor size and penalty from power supply companies in few countries. The above factors mention the importance of power factor improvement. Power factor correction can be proceeded by compensating the lagging current by connecting capacitor banks to the supply in small scale industries and homes. For household purposes, a capacitor with a capacitance of sufficient value is attached to bring the power factor almost to unity.

METHODOLOGY FOR POWER FACTOR CORRECTION SYSTEM

The methodology followed in this paper is presented in Figure 2. The capacitor bank is kept across the load and whenever the power factor adjustments are required the capacitors with proper value are put in parallel to the load. The values are converted into appropriate amounts for the Arduino UNO and phase delay measuring circuit using the current and voltage sensors. Zero Crossing Detector [ZCD] is used to measure the phase delay.





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The Arduino UNO receives this amount in order to determine the power factor. It creates a control signal to turn on the delay if the power factor is less than 0.9. Then, the relay gets activated and makes the capacitor to be connected with the load. Then the procedure is repeated until the power factor becomes greater than 0.9 and the proper capacitor might be Current sensor employed in this project is ACS712ELCTR-5B-T and provides precise current measurement up to 5A for both AC and DC signals. It is a fully integrated linear current sensor with low resistance current conductor and excellent voltage isolation based on the Hall Effect. [7-8]. The sensitivity of ACS712 ELCTR-5B-T is 185 mV/A. The output analog voltage is proportional to current to be measured. Even with loads running at high voltages, such as 230 V AC mains, the sensing terminal can detect current since the output sensed voltage is isolated from the measuring port.

The voltage sensor ZMPT101B can be used to measure the AC voltage up to 250 volts [9-10]. It is simple to operate and generates very accurate voltage and power. It has a multiturn trim potentiometer that may be used to calibrate the Arduino's ADC output. When there is no load connected, the sensor has an initial voltage (Offset) of 2.5V if the sensor is energised by 5 V DC. Ground loops and unintentional test equipment grounding are prevented by the isolation transformer in the voltage sensor, which separates it from the power line ground connection. Moreover, they reduce high-frequency noise that is generated by the power source. The ATMEGA328 microcontroller offers the board flexibility by allowing the controller chip to be changed or removed in the event that it becomes damaged or stops working properly. The Arduino UNO boards are the only ones with these flexible capabilities. ZCD is a simple comparator circuit with a ground line attached to one of the input pins, and a ZCD circuit schematic is displayed in the Figure 3.

The circuit is provided with zero voltage as reference to one of the inputs terminals to find zero crossing points of a sine wave. ZCD compares the input sinusoidal signal with the zero-voltage level. A digital high output pulse is produced each time a sine wave crosses from zero reference voltage to another value. The differential voltage is positive when the input grows from zero to positive, which causes the OP-AMP to attain positive saturation (+Vsat). The OP-AMP reaches the negative saturation voltage, or Vsat, when the input drops from zero to negative since the differential voltage is negative. It can be seen that the output switches between +Vsat and -Vsat whenever the input crosses zero level thus generating a square wave as output. An XOR gate has these two square waves attached to it as inputs. If there is any phase delay in the input, the XOR gate will be set to 1. The output of the XOR gate spikes with very little time delay when the load is resistive.

IMPLEMENTATION OF POWER FACTOR CORRECTION

The proposed method is simulated using Proteus simulator and then the hardware is implemented. The results are discussed in this section.

Simulation Results

Proteus is a potent simulation software tool and Engineers may correctly model the behavior of electronic circuits with this comprehensive suite. Proteus simulates the operation of the circuit in real time using sophisticated algorithms and models, enabling engineers to test and improve their ideas—before they are produced. The proposed circuit includes current transformer (CT), potential transformer [PT], ZCD circuit, Arduino UNO, capacitor and relay. The CT and PT sense the current and the voltage respectively and both the quantities—are fed into ZCD circuit to convert the output sinusoidal waveforms to square waveform. This data is fed to the Arduino—UNO in which the respective code to calculate the power factor is preloaded. The Figure 4 shows the simulation circuit for—automatic power factor correction circuit. In this circuit, the expected power factor is set as 0.95 and whenever the power factor of the load is less than 0.95—the relay—is activated which connects the capacitor across the load to compensate for the lagging power factor [11-15].

The Arduino code for power factor calculation based on the input derived from ZCD and relay activation code are written using Arduino IDE and uploaded in the Proteus software. The code used n this project is detailed in Figure 6. The IDE software utilizes an inductive counter to count the number of cycles where the input signal lags the output signal





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and a counter to keep track of the total number of cycles. The estimated power factor is displayed on the LCD. The setup function in this block of code, is called once the program launches. It establishes the input, output, and LED pin modes, initializes the display screen, initializes the serial communication. Also, it adds a delay and prints various notifications on the LCD. The primary portion of the program that executes repeatedly following the setup function is the loop function. The phase angle, counter, inductive counter, power factor, and a Boolean value for the power factor correction status are all initialized by these lines. To determine the phase angle and power factor for 50 cycles of the input signal, this block of code uses a while loop. The input voltage is first read using the analog Read function, followed by adding a delay and setting the output pin to HIGH. The phase angle is then determined using the cosine function, and it is then added to the angle variable. To count the number of cycles when the input signal lags behind the output signal, the software increments the counter and checks if the voltage is less than half of the maximum value (512). Eventually, another delay is added to finish one cycle, and the output pin is set to LOW.

HARDWARE RESULTS

The components are assembled and connected according to the simulation and the components used are given in the Table 1. The Figure 7 shows the hardware setup and Figure 8 shows the results. The ACS712ELECTR-6B-T is connected in series with 220 V supply. The V_{cc} pins of the sensor are connected to the 5V DC supply. The ZMPTB10 is also connected across the input supply. The output of the sensors is connected b the OPAMP as inputs through 10 k Ω resistors to form the ZCD. The offset voltage is reduced using the anti-parallel diodes across the OPAMP's input terminals. The sensor readings are processed on the Arduino board, and the necessary computations for power factor correction are done. The various voltage regulators IC7806 and IC7812 are combined with a bridge rectifier diode to suit the circuit's needs for a DC supply. The Arduino is programmed in such a way that when the power factor becomes less than 0.9 the output pin connected to the relay is set high. The relay is used to switch in a capacitor of 9 μ F to accurate the power factor and is linked in parallel with the load [16]. The power rating of the load and the necessary power factor correction are taken into consideration while choosing the capacitor. The load's power factor is improved and energy consumption is decreased after the power factor correction capacitor is turned on. The Arduino board keeps track of the load's voltage and currentwaveforms and makes the calculations required to keep the power factor at or near 1. Tekronix DSO is used to capture the output signal of sensors and ZCD and shown in Figure 9.

CONCLUSION

Energy monitoring innovations have benne grown during the past few years and effective strategies must be employed to enable effective, dependable utility management of diverse sectors. Hence a better and user-friendly method can be obtained by the enhancement of the conventional methods. Simple homes can be upgraded into smart homes with greater efficiency by implementing smart meter technology, appliance management and power factor correction methods. There are several potential future innovations that could improve the performance of smart homes including energy storage integration, use of advanced power electronics components, artificial intelligence-based control, and integration with smart grid technology.

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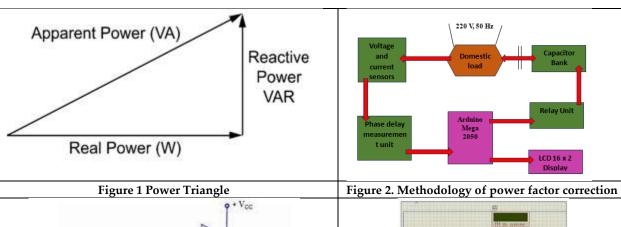
Table 1 List of components

Component	SPECIFICATIONS
Current sensor	ACS712ELECTR-6B-T
Voltage sensor	ZMPT101B
Arduino board	Arduino UNO
LCD	JHD204
XOR Gate	HEF4030B
OP AMP	TS324
Capacitor bank	9 micro farad
Motor load	40 W,220 V, 50 Hz





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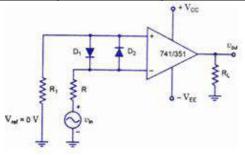


Figure 3. Circuit Diagram of Zero Crossing Detector

Figure 4. Automatic Power Factor Correction Unit

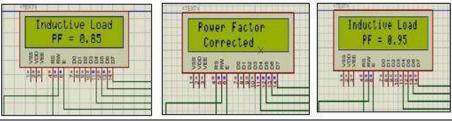


Figure 5. LCD display showing power factor before and after the correction

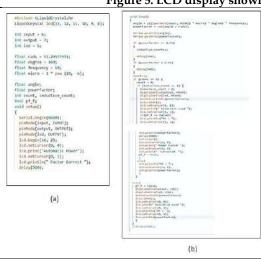




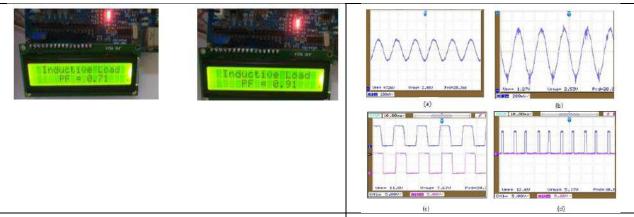
Figure 6. Arduino code for a) power factor calculation b) detecting lagging power factor and correcting the power factor

Figure 7 Hardware implementation of automatic power factor correction circuit





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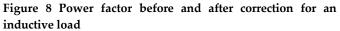


Figure 9. Output signals of (a) current sensor (b) voltage sensor (c) OPAMP (d) XOR gate of ZCD





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RESEARCH ARTICLE

Pilot Study on Concordance of Microsatellite Instability (MSI) Testing by Idylla Point of Care PCR with Immunohisto chemistry

Lavanya.N^{1*}, Sussela.K², Pratyusha.A³, Subramanyeshwar Rao.T⁴ and Raju. K.V.V.N⁵

Sr.Scientific Officer, Department of Molecular Pathalogy, Laboratory Medicine Basavatarakam Indo American Cancer Hospital & Research Institute (BIACH&RI), Hyderabad, Telangana, India.

²HoD, Department of Laboratory Medicine, Basavatarakam Indo-American Cancer Hospital and Research Institute, Hyderabad, Telangana, India.

³Junior Research Fellow, Department of Molecular Pathalogy, Laboratory Medicine, Basavatarakam Indo American Cancer Hospital & Research Institute (BIACH&RI), Hyderabad, Telangana, India.

⁴Medical Director, Surgical Oncologist, Department of Surgical oncology, Basavatarakam Indo American Cancer Hospital & Research Institute (BIACH&RI), Hyderabad, Telangana, India.

⁵Senior Consultant, Department of Surgical Oncology, Basavatarakam Indo American Cancer Hospital & Research Institute (BIACH&RI), Hyderabad, Telangana, India.

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*Address for Correspondence

Lavanya.N

Sr.Scientific Officer,

Department of Molecular Pathalogy,

Laboratory Medicine Basavatarakam Indo American Cancer Hospital & Research Institute (BIACH&RI), Hyderabad, Telangana, India.

Email: nambaru.lavanya@gmail.com



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ABSTRACT

Detection of microsatellite instability (MSI) is recommended for all patients with colorectal cancer (CRC). The prototype IdyllaTM MSI Test has been developed using a new set of short homopolymers located in the ACVR2A, BTBD7, DIDO1, MRE11, RYR3, SEC31A & SULF2 genes. This marker set allows probebased detection with great specificity in a simplified workflow compared to current methods. In this study, repeat length with this set of biomarkers was determined on formalin-fixed and paraffinembedded (FFPE) CRC samples using IdyllaTM MSI Test prototype cartridges, which allow a fully automated workflow including sample preparation, DNA amplification and automated repeat length calling. A total number of 18 samples including 14 adenocarcinomas from colorectal cancer (15 primary, one metastatic deposit in liver) and 4 metastatic deposits in liver including 2 from breast and one each from gallbladder and endometrium were included in the study. The grade of colorectal adenocarcinoma was grade 1 in 6, grade 2 in 7 and grade 3 in one. The viable tumor percentage ranged from 20-80%. Thirteen samples (13/18: 72 %) were microsatellite stable (MSS) by Idylla™ and five (5/18: 28 %) samples were MSI-H. Mutations were observed in ACVR 2A (5), DIDO1 (5), SULF2 (4), BTBD7 (4), RYR3 (3),





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SEC31A (2) and MRE11 (2) in the 5 MSI-H samples. Therefore, tests like these can work as effective and rapid diagnostic tools and provide personalized treatment for the patient. However, the limitation of the present study is the small number of samples and therefore sensitivity analysis could not be performed

Keywords: Idylla point, Immunohistochemistry, Microsatellite instability, PCR

INTRODUCTION

Microsatellites are short sequences of DNA (consisting of one to six bases) that repeat themselves tandemly, across the human genome. Since they constitute several repeats, they are susceptible to mismatch errors during DNA replication. If the DNA mismatch repair system is impaired due to mutations in genes involved, namely- mutL homologue 1 (MLH1), mutS homologue 2 (MSH2), mutS homologue 6 (MSH6) and post-meiotic segregation increased 2 (PMS2), it would result in the accumulation of such repeat length alterations, resulting in microsatellite instability (MSI) (1). The Food and Drug Adminstration (FDA) has granted accelerated approval to an anti-PD-1 antibody, pembrolizumab, for use in pediatric and adult patients with MSI+ solid tumors (3). MSI is therefore considered a potential predictive marker for chemotherapy and immunotherapy. Twenty percent of colorectal cancers (CRC) are found to be defective in mismatch repair (MMR) system while the remaining 80% are said to be caused due to chromosomal instability (3). Deficiency in the MMR system can also be associated with Lynch syndrome, a hereditary cancer syndrome that makes one susceptible to multiple types of cancer. Therefore, if a deficiency is detected, the patient can further be selected for a germline test in order to assess their status and manage their treatment accordingly (4). Mismatch repair deficiency/microsatellite instability-high connotes a good prognosis in early colorectal cancer settings without adjuvant treatment and a poor prognosis in patients harboring metastases(5). Their diagnosis can be done by molecular methods or by means of immunohistochemistry (IHC).

Molecular testing primarily involves profiling mono- and di-nucleotide(or high order nucleotides) loci. If all the loci are found to be stable, then it would result in MSS (microsatellite stable); if two or more loci show instability it would be called MSI-H (microsatellite instability- high) and if one locus shows instability it would be called MSI-L (microsatellite instability-low) (3). However, IHC involves the use of antibodies against the proteins encoded by the above-mentioned mismatch repair genes, MLH1, MSH2, MSH6 and PMS2. MLH1/PMS2 and MSH2/MSH6 occur as heterodimers and loss of either of them or even loss of expression of a single protein would imply impaired/ defective mismatch repair (1). Although IHC shows concordance with molecular testing; however, there are a few limitations. A non-functional protein caused due to missense mutations may be normally translated resulting in positive expression. On the other hand, it can also imply false positive results where-in, although there might have been loss of a protein expression, since it occurred only in a subset of mutated tumors, it might not affect the MMR machinery and still continue to remain efficient in DNA repair (1). MSI assay by IdyllaTM is a molecular techniquebased assay that uses real-time polymerase chain reaction (PCR) to assess the mutation status. It is an in vitro rapid diagnostic test that has been validated against Promega MSI analysis system and /or IHC in very few studies (2,7). Compared to Promega MSI Analysis System, an overall agreement, sensitivity and specificity of 98.7%, 94.4% and 100% have been reported with Idylla™ MSI Assay (2). In this study, we attempted to assess the concordance of MSI by IHC results with those obtained by using IdyllaTM MSI Assay (Biocartis, Belgium). It is a fully automated assay that processes the samples and gives an interpretation as well. It detects mutations in the following loci which as biomarkers are tumor specific and exhibit high frequency in colorectal and endometrial cancers: ACVR2A, BTBD7, DIDO1, MRE11, RYR3, SEC31A and SULF2. The present study is the first of its kind from India, to the best of our knowledge.





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MATERIALS AND METHODS

This was a prospective, observational study, approved by the institutional review board. Samples obtained for MSI IHC were included for validation. Informed consent was taken from all the patients. A total of 18 formalin fixed paraffin embedded (FFPE) tissue blocks were tested for MSI by both IHC and IdyllaTM. The site of biopsy and histology of each of the specimens were provided in table 1. One sample was tested for germline mutations in MLH1, MSH2, MSH6 and PMS2 on the Next Generation Sequencing (NGS) platform as well.

MSI Assay by IdyllaTM

Haematoxylin and eosin (H & E) stained slides of the tissue sections were reviewed by a pathologist. The protocol for sample processing was as per the protocol used by Lee M et al (6). A minimum of 20% viable tumor (VT) was taken as the cut off. Viable tumor percentage was ascertained by macro-dissection and tumor area was marked to determine the number of sections to be procured. The tissue blocks were refrigerated at 4°C for about 30 minutes prior to procuring the sections. The tissue was then processed in an Idylla™ instrument according to the manufacturer's instructions. The Biocartis Idylla™ MSI Assay was fully automated and capable of detecting seven monomorphic biomarkers. This qualitative detection technique uses FFPE tissues, from which nucleic acid is automatically extracted, PCRperformed and analyzed using high resolution melting detection using one cartridge alone.(in a single catridge). Specific PCR amplicons will generate biomarker specific fluorescence profiles. The validity of these profiles (which) is checked and analyzed by test specific software which(and) provides the final result to the user. Thus, this technique also eliminates the need for sample processing control.

Procedure for MMR testing by IHC

Procedure for IHC consisted of mounting of 4µm sections cut from FFPE blocks on poly -l-lysine coated slides, deparaffinized with xylene and hydrated through graded alcohols. The IHC was performed on BOND-III fully automated IHC stainer (Leica) as per manufacturer's instructions. The following antibodies were used: MLH1 (Clone GM011,DAKO, RTU), MSH2 (Clone FE11 DAKORTU)MSH6(Clone EP49 RTU,DAKO) PMS2(Clone EP51,DAKORTU) Every section of colon had normal mucosa, which served as internal positive control. However where internal positive controls were not available (eg metastatic sites small biopsy samples etc) external controls were also run. Samples with intact expression were considered to have intact MMR function and loss of expression of paired markers MLH1 and PMS2 or MSH2 and MSH6 were suggestive of MMR deficient status. Sole loss of any of these markers was reported as such and necessitated further testing including germline testing.

Procedure for MSI testing by NGS

An informed consent was taken for the patient for germline testing with hereditary cancer genes. Germline testing for hereditary cancer panel (including MLH1, MSH2, MSH6 and PMS2) was performed by NGS based assay using 30 genes and associated Bioinformatics pipeline.which showed pathogenic mutations in MSH2. The sequencing was performed on Ion TorrentS5 plus NGS platform at 400X coverage. Results were evaluated on the Ion Torrent software

RESULTS

A total number of 18 samples including 14 adenocarcinomas from colorectal cancer (15 primary, one metastatic deposit in liver) and 4 metastatic deposits in liver including 2 from breast and one each from gallbladder and endometrium were included in the study. The grade of colorectal adenocarcinoma was grade 1 in 6, grade 2 in 7 and grade 3 in one.

MSI assay by IdyllaTM and MMR status by IHC





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The viable tumor percentage ranged from 20-80%. Thirteen samples (13/18:72 %)were microsatellite stable (MSS) by IdyllaTM and five (5/18: 28 %) samples were MSI-H. Mutations were observed in ACVR 2A (5), DIDO1 (5), SULF2 (4), BTBD7 (4), RYR3 (3), SEC31A (2) and MRE11 (2) in the 5 MSI-H samples (Table 1). MMR status by IHC showed loss of expression of MLH1& PMS2 in 3 and MSH2 &MSH6 in two samples (Table 1). There was complete concordance between the results of IdyllaTM and IHC in the 18 samples tested. One patient who had adetailed work-up whose sample was subjected to IHC, Idylla and NGS is described below (Table 3- Case 18). A 40 years old patient who presented with adenocarcinoma of the transverse colon was initially evaluated by IHC for MMR status. There was loss of MSH2 and MSH6 expression and intact MLH1 and PMS2 expression. This was indicative of MSI -H status and therefore genomic testing by PCR was done by IdyllaTM, which demonstrated MSI-H status (all 7 biomarkers were mutant: MSI-H;mutation detected in all seven biomarkers ACVR2A, BTBD7, DIDO1, MRE11, RYR3, SEC31A and SULF2.) Thereafter, in view of high risk, germline testing for MSH2 and other genes implicated in Lynch Syndrome was performed for hereditary colon cancer panel (comprising of 30 genes) on Ion Torrent S5 plus NGS platform with 400X coverage, after obtaining informed consent. The test yielded a pathogenic mutation in MSH2. The changes were observed in chromosome 2, exon 5(Indel). The amino acid change was Leu280Phe.The patient was counselled and advised for testing of the other family members as well, to identify Lynch syndrome. This case illustrates the importance of the three different technologies.

DISCUSSION

The present study compared the results of MSI testing by IdyllaTM, that detects mutations with high frequency in colorectal and endometrial cancers, to the results obtained by IHC for MMR protein expression in 18 samples. There was 100% concordance of the results. Tissues, that showed loss of MLH1 and PMS2 markers by IHC, also showed mutations by IdyllaTM in the 3 biomarkers in common, namely ACVR2A, DIDO1 and SULF2. The former two mutations were found in tissue that showed loss of MSH2 and MSH6 as well (Table 2). All the tissues that were MSI-intact when detected by IHC were 100% concordant with the results of IdyllaTM which were microsatellite stable. The tumor tissues tested were from different regions of the colon (ascending, sigmoid, transverse colon), rectum and the liver. However, all the four tissues that showed MSI-H status belonged to different parts of the colon. The minimum viable tumor percentage that allowed mutation detection for these set of samples in IdyllaTM was 20% (Table 1). When the area of the tissue section with viable tumor was small, we obtained additional number of sections. IdyllaTM produces the result in approximately two and a half hours, whereas IHC would require several hours to overnight staining procedure. Furthermore, IdyllaTM detects mutations at the RNA (DNA) level which is more reliable than a protein-based test such as IHC. Moreover, IdyllaTM Assay does not require a control tissue unlike IHC.

Each of the five cases with loss of MMR expression and corresponding MSI-H status had a different presentation. Other alternatives to MSI status testing include real time PCR, Gene sequencing by fragment analysis and NGS. Real time PCR involves more steps as it requires extraction of the nucleic acid from the tissue, followed by its purification, quantification, PCR runs and analysis. NGS although provides a more accurate result, is time consuming, requires more consumables, instruments and also expensive. However, as in the illustrated case, NGS facilitated identification of a pathogenic mutation in MSH2, thus enabling the patient to be counselled and advised for testing the other family members as well to identify Lynch syndrome. The advantages of Idylla™ Assay include that console occupies less space, is easy to process and is quicker but the cartridge provided is only for a single use and is expensive as well. The lack of manual intervention once the tissue is inserted into cartridge can pose an advantage as well as a disadvantage. If the amount of tissue inserted or nucleic acid extracted is insufficient or any such extraction/processing error occurs, one would be able to view it only at the very end of the procedure and would have to use another cartridge and cut more sections of tissue to repeat the test. Although Idylla™ provides a QC report at the end of the test; one cannot view the amplification curve to assess the quality of tissue or nucleic acid.





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The set of mutations that IdyllaTM detects are of considerable importance. Although rare, false positive results have been reported where in despite efficient MMR machinery, somatic mutations were detected in 4 of the genes(6). Apart from validation of MSI results obtained through IdyllaTM with that of IHC, we have also found common mutations in tumors with a loss of MLH1 and PMS2. This could possibly pave way for establishing a gene signature that can further help in the better assessment of the pathological status of the tumor and also for an effective targeted therapy. Clinical findings may not be predictive of the results of biomarker testing. Therefore, tests like these can work as effective and rapid diagnostic tools and provide personalized treatment for the patient. However, the limitation of the present study is the small number of samples and therefore sensitivity analysis could not be performed in a laboratory setting, simple, easy, less time-consuming techniques are imperative considering the needs of the patient and the urgency for therapeutic planning by the clinician. Keeping these factors in mind, we believe IdyllaTM is a robust technology and good alternative in the clinic for rapid MSI testing.

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Table 1: Results of MSI on Idylla™ Assay and MMR status on IHC, site of biopsy and histology

Sample No.	Results on Idylla TM	ViableTumor Percentage	Result by IHC	Site of Biopsy	Histology	Grade
1	MSI-H	65%	MMR expression	Colon	Adenocarcinoma	3
2	MSI-H	70%	MMR	Transverse colon	Adenocarcinoma	1
3	MSS	20%	MMR	Rectum	Adenocarcinoma	2
4	MSS	80%	MMR	Rectosigmoid	Adenocarcinoma	1
5	MSS	70%	MMR	Rectum and	Adenocarcinoma	2





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6	MSS	60%	MMR	Rectum	Adenocarcinoma	1
7	MSI-H	60%	MMR	Sigmoid colon	Adenocarcinoma	1
8	MSS	20%	MMR expression	Liver (trucut biopsy)	Metastatic deposits of carcinoma endometrium.	
9	MSS	60%	MMR expression	Liver (trucut biopsy)	metastatic adenocarcinoma from	
10	MSS	50%	MMR expression	Liver (trucut biopsy)	Metastases from breast	
11	MSS	60%	MMR	Transverse colon	Adenocarcinoma	2
12	MSS	20%	MMR	Rectum	Adenocarcinoma	2
13	MSS	70%	MMR	Rectum	Adenocarcinoma	2
14	MSS	60%	MMR	Colon	Adenocarcinoma	2
15	MSS	40%	MMR	Liver	Metastatic	2
16	MSS	60%	MMR	Liver	Metastases from breast	
17	MSI-H		MMR	Ascending colon	Adenocarcinoma	1
18	MSI -H	40%	MMR	Transverse colon	Adenocarcinoma	1

Abbreviations: IHC: MMR: Mismatch repair; MSI-H: Microsatellite Instability High; MSS:-Microsatellite Stable

Table 2: Table demonstrating MSI-H status and IHC (n=5)

	Ic	dylla™	Bioma	arker S	Status-	(MSI-F	H)	IHC marker Status				
Sample No.	ACVR2A	DID01	SULF2	BTBD7	RYR 3	SEC 31A	MRE11	MLH1	PMS2	MSH2	MSH6	
1	+	+	+	+	+	+	-	Lost	Lost	Intact	Intact	
7	+	+	+	-	-	-	+	Lost	Lost	Intact	Intact	
17	+	+	+	+	-	-	-	Lost	Lost	Intact	Intact	
2	+	+	-	+	+	-	-	Intact	Intact	Lost	Lost	
18*	+	+	+	+	+	+	+	Intact	Intact	Lost	Lost	

⁺ Detected; - Not detected





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RESEARCH ARTICLE

On $\alpha^* g^{\dagger} \psi$ -closed Sets in Topological Spaces

S.Pooja^{1*}, M.Vigneshwaran² and L.Vidyarani ²

¹Ph.D Research Scholar, Department of Mathematics, Kongunadu Arts and Science College (Affiliated to Bharathiar University) Coimbatore, Tamil Nadu, India.

²Assistant Professor, Department of Mathematics, Kongunadu Arts and Science College (Affiliated to Bharathiar University) Coimbatore, Tamil Nadu, India.

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*Address for Correspondence

S.Pooja

Ph.D Research Scholar,

Department of Mathematics,

Kongunadu Arts and Science College (Affiliated to Bharathiar University)

Coimbatore, Tamil Nadu, India. Email: poojabreeks@gmail.com



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ABSTRACT

The subject of this work is Alpha star generalized hash psi closed $set(\alpha^* q^{\#} \psi)$ in topological spaces where the relation of a $\alpha^*g^{\#}\psi$ -closed set to other generalised sets are derived. Also the characteristics of $\alpha^*g^{\#}\psi$ closed set is derived.

Keywords: $\alpha^* g^{\#} \psi$ -closed set.

MSC: 54A05

INTRODUCTION

Levine [7, 8] developed generalised closed sets and semi- open sets in topological spaces. Njastad [14] introduced α sets. S.P.Arya and T.Nour[1] introduced gs-closed sets. H.Maki et al.[9,10] developed $g\alpha$ -closed sets and α g-closed sets. M.Vigneshwaran and R.Devi[18] developed the idea of *gα-closed sets. gsp-closed sets and gpr-closed sets were developed by Dontchev[3] and Gnanambal[4] respectively. Veerakumar[16] introduced ψ -closed sets. Kanimozhi, Balamani and Parvathi[6] introduced $g^{\#}\psi$ -closed sets and T.Nandhini[13] introduced $\alpha g^{\#}\psi$ -closed sets. The objective of this article is to develop a new group of sets named Alpha star generalized hash psi closed sets in topological spaces.

Preliminaries

Definition 2.1. Let P be a topological space, $S \subseteq T$ is stated as $sOS [8] if S\subseteq cl(int(S)) and <math>sCS if int(cl(S)) \subseteq S$.





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pOS [12] if $S\subseteq int(cl(S))$ and pCS if $cl(int(S))\subseteq S$. α OS [14] if S \subseteq int(cl(int(S))) and α CS if cl(int(cl(S))) \subseteq S. gCS [7] if $cl(S) \subseteq T$ when $S \subseteq T$ and T is open in P. gsCS [1] if $scl(S) \subseteq T$ when $S \subseteq T$ and T is open in P. gpCS [11] if $pcl(S) \subseteq T$ when $S \subseteq T$ and T is open in P. $g\alpha CS$ [9] if $\alpha cl(S) \subseteq T$ when $S\subseteq T$ and T is α -open in P. *g α CS [18] if cl(S) ⊆T when S⊆T and T is g α -open in P. sgCS [2] if $scl(S) \subseteq T$ when $S\subseteq T$ and T is semi open in P. α gCS [10] if α cl(S) \subseteq T when S \subseteq T and T is open in P. gspCS [3] if spcl(S) \subseteq T when S \subseteq T and T is open in P. g*spCS [5] if spcl(S) ⊆T when S⊆T and T is g-open in P. gprCS [4] if $pcl(S) \subseteq T$ when $S \subseteq T$ and T is regular open in P. ψ CS [16] if scl(S) ⊆T when S⊆T and T is sg-open in P. ψ gCS [15] if ψ cl(S) \subseteq T when S \subseteq T and T is open in P. $g^{\#}CS$ [17] if cl(S) $\subseteq T$ when $S\subseteq T$ and T is αg -open in P. $g^{\#}\psi$ CS [6] if ψ cl(S) \subseteq T when S \subseteq T and T is ψ -open in P. $\alpha g^{\#}\psi$ CS [13] if α cl(S) \subseteq T when S \subseteq T and T is $g^{\#}\psi$ -open in P.

Basic Properties of $\alpha^* g^\# \psi$ -Closed Set

Definition 3.1: Let P be a topological space, a subset S of P is stated as $\alpha^* g^\# \psi$ -closed set($\alpha^* g^\# \psi$ -CS) if α cl(S) \subseteq T when S \subseteq T and T is $\alpha g^\# \psi$ -open set($\alpha g^\# \psi$ -OS).

Theorem 3.2: Every closed set is $\alpha^* g^{\#} \psi$ -CS.

Proof: Let $S \subseteq T$, T is $\alpha g^{\#} \psi$ -OS in P. As S is closed, cl(S) = S. But $\alpha cl(S) \subseteq cl(S)$. $\Rightarrow \alpha cl(S) \subseteq T$. Hence S is $\alpha^* g^{\#} \psi$ -CS.

The below example demonstrates the preceding implication is irreversible.

```
Example 3.3: Let P = \{x_1, x_2, x_3\}, \tau = \{P, \emptyset, \{x_1, x_3\}\}. \alpha^* g^\# \psi C(P, \tau) = \{P, \emptyset, \{x_2\}, \{x_1, x_2\}, \{x_2, x_3\}\}. \{x_1, x_2\} is \alpha^* g^\# \psi-CS of (P, \tau) although not closed in (P, \tau).
```

Theorem 3.4: Every α -CS is $\alpha^* g^\# \psi$ -CS. **Proof:** Let S \subseteq T, T is $\alpha g^\# \psi$ -OS in P. As S is α -CS, α cl(S) = S. But α cl(S) \subseteq T. Hence S is $\alpha^* g^\# \psi$ -CS.

The below example demonstrates the preceding implication is irreversible.

```
Example 3.5: Let P = \{x_1, x_2, x_3\}, \tau = \{P, \emptyset, \{x_2, x_3\}\}.

\alpha C(P, \tau) = \{P, \emptyset, \{x_1\}\}.

\alpha^* g^\# \psi C(P, \tau) = \{P, \emptyset, \{x_1\}, \{x_1, x_2\}, \{x_1, x_3\}\}.

\{x_1, x_2\} is \alpha^* g^\# \psi -CS of (P, \tau) although not \alpha-CS in (P, \tau).
```

Theorem 3.6: Every $\alpha^* g^\# \psi$ -CS is gs-CS. **Proof:** Let S \subseteq T, T is OS in P. As every OS is $\alpha g^\# \psi$ -OS, T is $\alpha g^\# \psi$ -OS.





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Here S is $\alpha^* g^\# \psi$ -CS,

 $\Rightarrow \alpha cl(S) \subseteq T$.

Also $scl(S) \subseteq \alpha cl(S)$, then $scl(S) \subseteq T$.

Hence S is gs-CS.

The below example demonstrates the preceding implication is irreversible.

Example 3.7: Let $P = \{x_1, x_2, x_3\}$, $\tau = \{P, \emptyset, \{x_2\}, \{x_1, x_2\}\}$. GSC(P, τ) = $\{P, \emptyset, \{x_1\}, \{x_3\}, \{x_2, x_3\}, \{x_1, x_3\}\}$. $\alpha^* g^\# \psi C(P, \tau) = \{P, \emptyset, \{x_1\}, \{x_3\}, \{x_1, x_3\}\}$. $\{x_2, x_3\}$ is gs-CS of (P, τ) although not $\alpha^* g^\# \psi$ -CS in (P, τ).

Theorem 3.8: Every $\alpha^* g^{\#} \psi$ -CS is gp-CS.

Proof: Let $S\subseteq T$, T is OS in P.

As every OS is $\alpha g^{\#}\psi$ -OS, T is $\alpha g^{\#}\psi$ -OS.

Here S is $\alpha^* g^{\#} \psi$ -CS.

 $\Rightarrow \alpha cl(S) \subseteq T$.

Also, $pcl(S) \subseteq cl(S)$.

 \Rightarrow pcl(S) \subseteq T.

Hence S is gp-CS.

The below example demonstrates the preceding implication is irreversible.

Example 3.9: Let $P = \{x_1, x_2, x_3\}$, $\tau = \{P, \emptyset, \{x_2, x_3\}\}$. GPC(P, τ) = $\{P, \emptyset, \{x_1\}, \{x_2\}, \{x_3\}, \{x_1, x_2\}, \{x_1, x_3\}\}$. $\alpha^* g^\# \psi C(P, \tau) = \{P, \emptyset, \{x_1\}, \{x_1, x_2\}, \{x_1, x_3\}\}$. $\{x_2\}$ is gp-CS of (P, τ) although not $\alpha^* g^\# \psi$ -CS in (P, τ).

Theorem 3.10: Every $\alpha^* g^{\#} \psi$ -CS is sg-CS.

Proof: Let $S \subseteq T$, T is semi-OS in P.

As every semi-OS is $\alpha g^{\#}\psi$ -OS, T is $\alpha g^{\#}\psi$ -OS.

As S is $\alpha^* g^\# \psi$ -CS, α cl(S) \subseteq T.

Also $scl(S) \subseteq \alpha cl(S)$.

Then, $scl(S) \subseteq T$.

Hence S is sg-CS.

The below example demonstrates the preceding implication is irreversible.

Example 3.11: Let $P = \{x_1, x_2, x_3\}$, $\tau = \{P, \emptyset, , \{x_1\}, \{x_2\}, \{x_1, x_2\}\}$. SGC(P, τ) = $\{P, \emptyset, \{x_1\}, \{x_2\}, \{x_3\}, \{x_2, x_3\}, \{x_1, x_3\}\}$. $\alpha^* g^\# \psi C(P, \tau) = \{P, \emptyset, \{x_3\}, \{x_2, x_3\}, \{x_1, x_3\}\}$. $\{x_1\}$ is sg-CS of (P, τ) although not $\alpha^* g^\# \psi$ -CS in (P, τ).

Theorem 3.12: Every $\alpha^* g^\# \psi$ -CS is αg -CS.

Proof: Let $S \subseteq T$, T is OS in P.

As every OS is $\alpha g^{\#}\psi$ -OS, T is $\alpha g^{\#}\psi$ -OS.

Here S is $\alpha^* g^{\#} \psi$ -CS.

 $\Rightarrow \alpha cl(S) \subseteq T$.

Hence S is α g-CS.

The below example demonstrates the preceding implication is irreversible.





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Example 3.13: Let $P = \{x_1, x_2, x_3\}$, $\tau = \{P, \emptyset, \{x_3\}, \{x_2, x_3\}\}$. $\alpha GC(P, \tau) = \{P, \emptyset, \{x_1\}, \{x_2\}, \{x_1, x_2\}, \{x_1, x_3\}\}$. $\alpha^* g^\# \psi C(P, \tau) = \{P, \emptyset, \{x_1\}, \{x_2\}, \{x_1, x_2\}\}$. $\{x_1, x_3\}$ is αg -CS of (P, τ) although not $\alpha^* g^\# \psi$ -CS in (P, τ) .

Theorem 3.14: Every * $g\alpha$ -CS is α *g# ψ -CS.

Proof: Let $S \subseteq T$, T is $\alpha g^{\#} \psi$ -OS in P.

As every open set T is $\alpha g^{\#}\psi$ -OS, T is $\alpha g^{\#}\psi$ -OS.

Since S is *g α -CS. Then, cl(S)⊆T.

Here, $\alpha cl(S) \subseteq cl(S)$.

 $\Rightarrow \alpha \operatorname{cl}(S) \subseteq T$.

Hence S is $\alpha^* g^\# \psi$ -CS.

The below example demonstrates the preceding implication is irreversible.

Example 3.15: Let $P = \{x_1, x_2, x_3\}, \tau = \{P, \emptyset, \{x_1\}\}.$ * $G\alpha C(P, \tau) = \{P, \emptyset, \{x_2, x_3\}\}.$ $\alpha^* g^\# \psi C(P, \tau) = \{P, \emptyset, \{x_2\}, \{x_3\}, \{x_2, x_3\}\}.$ $\{x_2\}$ is $\alpha^* g^\# \psi$ -CS of (P, τ) although not * $g\alpha$ -CS in (P, τ) .

Theorem 3.16: Every $\alpha^* g^\# \psi$ -CS is gsp-CS.

Proof: Let $S\subseteq T$, T is OS in P.

As every open set T is $\alpha g^{\#}\psi$ -OS, T is $\alpha g^{\#}\psi$ -OS.

Since S is $\alpha^* g^\# \psi$ -CS, α cl(S) \subseteq T.

Also $\operatorname{spcl}(S) \subseteq \alpha \operatorname{cl}(S)$.

 \Longrightarrow spcl(S) \subseteq T.

Hence S is gsp-CS.

The below example demonstrates the preceding implication is irreversible.

Example 3.17: Let $P = \{x_1, x_2, x_3\}, \tau = \{P, \emptyset, \{x_1, x_2\}\}.$ GSPC(P, τ) = $\{P, \emptyset, \{x_1\}, \{x_2\}, \{x_3\}, \{x_2, x_3\}, \{x_1, x_3\}\}.$ $\alpha^* g^\# \psi C(P, \tau) = \{P, \emptyset, \{x_3\}, \{x_2, x_3\}, \{x_1, x_3\}\}.$ $\{x_1\}$ is gsp-CS of (P, τ) although not $\alpha^* g^\# \psi$ -CS in (P, τ).

Theorem 3.18: Every $\alpha^* g^\# \psi$ -CS is g*sp-CS.

Proof: Let $S\subseteq T$, T is g-OS in P.

As every g-OS is $\alpha g^{\#}\psi$ -OS, T is $\alpha g^{\#}\psi$ -OS.

Since S is $\alpha^* g^\# \psi$ -CS, α cl(S) \subseteq T.

But $spcl(S) \subseteq \alpha cl(S)$.

 \Rightarrow spcl(S) \subseteq T.

Hence S is g*sp-CS.

The below example demonstrates the preceding implication is irreversible.

Example 3.19: Let $P = \{x_1, x_2, x_3\}, \tau = \{P, \emptyset, \{x_1\}, \{x_1, x_2\}\}\}$. G*SPC(P, τ) = $\{P, \emptyset, \{x_2\}, \{x_3\}, \{x_2, x_3\}, \{x_1, x_3\}\}$. $\alpha^* g^\# \psi C(P, \tau) = \{P, \emptyset, \{x_2\}, \{x_3\}, \{x_2, x_3\}\}$. $\{x_1, x_3\}$ is g*sp-CS of (P, τ) although not $\alpha^* g^\# \psi$ -CS in (P, τ).

Theorem 3.20: Every $\alpha^* g^\# \psi$ -CS is gpr-CS.

Proof: Let $S\subseteq T$, T is regular OS in P.





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As every regular OS is $\alpha g^{\#}\psi$ -OS, T is $\alpha g^{\#}\psi$ -OS.

Since S is $\alpha^* g^\# \psi$ -CS, α cl(S) \subseteq T.

Also $pcl(S) \subseteq \alpha cl(S)$.

 \Rightarrow pcl(S) \subseteq T.

Hence S is gpr-CS.

The below example demonstrates the preceding implication is irreversible.

Example 3.21: Let $P = \{x_1, x_2, x_3\}$, $\tau = \{P, \emptyset, \{x_2\}\}$. GPRC(P, τ) = $\{P, \emptyset, \{x_1\}, \{x_2\}, \{x_3\}, \{x_1, x_2\}, \{x_2, x_3\}, \{x_1, x_3\}\}$. $\alpha^* g^\# \psi C(P, \tau) = \{P, \emptyset, \{x_1\}, \{x_3\}, \{x_1, x_3\}\}$. $\{x_2\}$ is gpr-CS of (P, τ) although not $\alpha^* g^\# \psi$ -CS in (P, τ).

Theorem 3.22: Every $\alpha^* g^\# \psi$ -CS is ψ g-CS.

Proof: Let $S \subseteq T$, T is OS in P.

As every OS is $\alpha g^{\#}\psi$ -OS, T is $\alpha g^{\#}\psi$ -OS.

Since S is $\alpha^* g^\# \psi$ -CS, α cl(S) \subseteq T.

Also ψ cl(S) $\subseteq \alpha$ cl(S).

 $\Rightarrow \psi cl(S) \subseteq T$.

Hence S is ψ g-CS.

The below example demonstrates the preceding implication is irreversible.

Example 3.23: Let $P = \{x_1, x_2, x_3\}$, $\tau = \{P, \emptyset, \{x_1\}, \{x_1, x_2\}\}$. ψ GC(P, τ) = $\{P, \emptyset, \{x_2\}, \{x_3\}, \{x_2, x_3\}, \{x_1, x_3\}\}$. $\alpha^* g^\# \psi$ C(P, τ) = $\{P, \emptyset, \{x_2\}, \{x_3\}, \{x_2, x_3\}\}$. $\{x_1, x_3\}$ is ψ g-CS of (P, τ) although not $\alpha^* g^\# \psi$ -CS in (P, τ).

Theorem 3.24: Every $g^{\#}$ -CS is $\alpha^*g^{\#}\psi$ -CS.

Proof: Let $S\subseteq T$, T is $\alpha g^{\#}\psi$ -OS in P.

As S is $g^\#$ -CS, then cl(S) ⊆T.

But $\alpha cl(S) \subseteq cl(S)$.

 $\Rightarrow \alpha cl(S) \subseteq T$.

Hence S is $\alpha^* g^\# \psi$ -CS.

The below example demonstrates the preceding implication is irreversible.

Example 3.25: Let $P = \{x_1, x_2, x_3\}, \tau = \{P, \emptyset, \{x_3\}, \{x_1, x_2\}\}.$ $g^{\#}C(P, \tau) = \{P, \emptyset, \{x_3\}, \{x_1, x_2\}\}.$ $\alpha^*g^{\#}\psi C(P, \tau) = \{P, \emptyset, \{x_1\}, \{x_2\}, \{x_3\}, \{x_1, x_2\}, \{x_2, x_3\}, \{x_1, x_3\}\}.$ $\{x_1\}$ is $\alpha^*g^{\#}\psi$ -CS of (P, τ) although not $g^{\#}$ -CS in (P, τ) .

Theorem 3.26: Every $\alpha^* g^\# \psi$ -CS is $g^\# \psi$ -CS.

Proof: Let $S \subseteq T$, T is ψ -OS in P.

As every ψ -OS is $\alpha g^{\#}\psi$ -OS, T is $\alpha g^{\#}\psi$ -OS.

As S is $\alpha^* g^\# \psi$ -CS, α cl(S) \subseteq T.

But ψ cl(S) $\subseteq \alpha$ cl(S).

 $\Rightarrow \psi cl(S) \subseteq T$.

Hence S is $g^{\#}\psi$ -CS.

The below example demonstrates the preceding implication is irreversible.

Example 3.27: Let $P = \{x_1, x_2, x_3\}, \tau = \{P, \emptyset, \{x_1\}, \{x_2\}, \{x_1, x_2\}\}.$





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```
g^{\#}\psi\mathsf{C}(\mathsf{P},\tau) = \{P,\emptyset,\{x_1\},\ \{x_2\},\ \{x_3\},\ \{x_2,x_3\},\ \{x_1,x_3\}\}. \alpha^*g^{\#}\psi\mathsf{C}(\mathsf{P},\tau) = \{P,\emptyset,\{x_3\},\ \{x_2,x_3\},\ \{x_1,x_3\}\} \{x_1\} \text{ is } g^{\#}\psi\mathsf{-CS} \text{ of } (\mathsf{P},\tau) \text{ although not } \alpha^*g^{\#}\psi\mathsf{-CS} \text{ in } (\mathsf{P},\tau).
```

Theorem 3.28: Every $\alpha g^{\#}\psi$ -CS is $\alpha^*g^{\#}\psi$ -CS.

Proof: Let $S\subseteq T$, T is $\alpha g^{\#}\psi$ -OS in P.

As every $\alpha g^{\#}\psi$ -OS is $g^{\#}\psi$ -OS, T is $g^{\#}\psi$ -OS.

As S is $\alpha g^{\#}\psi$ -CS, α cl(S) \subseteq T.

Hence S is $\alpha^* g^* \psi$ -CS.

The below example demonstrates the preceding implication is irreversible.

```
Example 3.29: Let P = \{x_1, x_2, x_3\}, \tau = \{P, \emptyset, \{x_2\}, \{x_1, x_3\}\}. \alpha g^{\#}\psi C(P, \tau) = \{P, \emptyset, \{x_2\}, \{x_1, x_3\}\}. \alpha^* g^{\#}\psi C(P, \tau) = \{P, \emptyset, \{x_1\}, \{x_2\}, \{x_3\}, \{x_1, x_2\}, \{x_2, x_3\}, \{x_1, x_3\}\}. \{x_3\} is \alpha^* g^{\#}\psi - CS of \{P, \tau\} although not \alpha g^{\#}\psi - CS in \{P, \tau\}.
```

Characteristics of $\alpha^* g^{\#} \psi$ -Closed Set

Remark 4.1: $\alpha^* g^\# \psi$ -CS is independent of semi-CS and ψ -CS.

It can be shown by the following examples.

Example 4.2: Let $P = \{x_1, x_2, x_3\}, \tau = \{P, \emptyset, \{x_1\}, \{x_2\}, \{x_1, x_2\}\}.$

SC(P, τ) = {P, \emptyset , { x_1 }, { x_2 }, { x_3 }, { x_2 , x_3 }, { x_1 , x_3 } = ψ C(P, τ).

 $\alpha^* g^\# \psi C(P, \tau) = \{P, \emptyset, \{x_3\}, \{x_2, x_3\}, \{x_1, x_3\}\}.$

Here $\{x_2\}$ is semi-CS and ψ -CS of (P, τ) although not $\alpha^* g^\# \psi$ -CS in (P, τ) .

Let P = $\{x_1, x_2, x_3\}, \tau = \{P, \emptyset, \{x_3\}, \{x_1, x_2\}\}.$

 $SC(P, \tau) = \{P, \emptyset, \{x_3\}, \{x_1, x_2\}\}. = \psi C(P, \tau).$

 $\alpha^* g^{\#} \psi C(P, \tau) = \{P, \emptyset, \{x_1\}, \{x_2\}, \{x_3\}, \{x_1, x_2\}, \{x_2, x_3\}, \{x_1, x_3\}\}.$

Here $\{x_1\}$ is $\alpha^* g^\# \psi$ -CS of (P, τ) although not semi-CS and ψ -CS in (P, τ) .

Theorem 4.3: The intersection of two $\alpha^* g^\# \psi$ -CS is again $\alpha^* g^\# \psi$ -CS.

Proof: Let X and Y be $\alpha^* g^\# \psi$ -CS and X\(\Gamma\)Y\(\subseteq\T\) where T is $\alpha g^\# \psi$ -OS.

As X and Y are $\alpha^* g^\# \psi$ -CS, $\alpha cl(X) \subseteq T$ and $\alpha cl(Y) \subseteq T$.

Also $\alpha \operatorname{cl}(X \cap Y) = \alpha \operatorname{cl}(X) \cap \alpha \operatorname{cl}(Y) \subseteq T$.

 $\Rightarrow \alpha \operatorname{cl}(X \cap Y) \subseteq T$ where $X \cap Y$ is $\alpha^* g^\# \psi$ -CS.

Hence, the intersection of two $\alpha^* g^\# \psi$ -CS is again $\alpha^* g^\# \psi$ -CS.

Theorem 4.4: The union of two $\alpha^* g^\# \psi$ -CS is again $\alpha^* g^\# \psi$ -CS.

Proof: Let X and Y be $\alpha^* g^\# \psi$ -CS and XUY \subseteq T where T is $\alpha g^\# \psi$ -OS.

As X and Y are $\alpha^* g^{\#} \psi$ -CS, $\alpha cl(X) \subseteq T$ and $\alpha cl(Y) \subseteq T$.

Also $\alpha \operatorname{cl}(X \cup Y) = \alpha \operatorname{cl}(X) \cup \alpha \operatorname{cl}(Y) \subseteq T$.

 $\Rightarrow \alpha \operatorname{cl}(X \cup Y) \subseteq T$ where XUY is $\alpha^* g^\# \psi$ -CS.

Hence, the union of two $\alpha^* g^\# \psi$ -CS is again $\alpha^* g^\# \psi$ -CS.

Theorem 4.5: Let X be an OS and Y be an $\alpha^* g^{\#} \psi$ -OS, then XUY is $\alpha^* g^{\#} \psi$ -OS.

Proof: Let X be an OS of (P, τ) and Y be an $\alpha^* g^{\#} \psi$ -OS of (P, τ)

As every OS is $\alpha^* g^\# \psi$ -OS, X is $\alpha^* g^\# \psi$ -OS.

Hence XUY is $\alpha^* g^\# \psi$ -OS.

Since union of two $\alpha^* g^\# \psi$ -OS is again $\alpha^* g^\# \psi$ -OS.

Theorem 4.6: Let X be an $\alpha^* g^\# \psi$ -CS of (P, τ) iff α cl(X)- X $\not\subseteq$ any non-empty $\alpha g^\# \psi$ -CS.





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Proof:

Necessary Part: Assume X is $\alpha^* g^\# \psi$ -CS and F be a non empty $\alpha g^\# \psi$ -CS with $F \subseteq \alpha cl(X)$ - X.

Then $X \subseteq P$ - F

 $\Rightarrow \alpha \operatorname{cl}(X) \subseteq P - F$.

Hence, $F \subseteq P$ - $\alpha \operatorname{cl}(X)$, which contradicts.

Sufficient Part : Assume X is a subset of (P, τ) such that $\alpha cl(X)$ - X $\not\subseteq$ any non- empty $\alpha g^{\#}\psi$ -CS.

Let T be an $\alpha g^{\#}\psi$ -OS in (P, τ) such that X \subseteq T.

If $\alpha \operatorname{cl}(X) \subseteq T$, then $\alpha \operatorname{cl}(X) \cap \operatorname{C}(T) \neq \emptyset$.

Then $\emptyset \neq \alpha \operatorname{cl}(X) \cap C(T)$ is an $\alpha g^{\#}\psi$ -CS of (P, τ) , since the intersection of two $\alpha g^{\#}\psi$ -CS is again $\alpha g^{\#}\psi$ -CS.

Theorem 4.7: If X is $\alpha^* g^\# \psi$ -CS and X \subseteq Y $\subseteq \alpha$ cl(X), then Y is $\alpha^* g^\# \psi$ -CS.

Proof: Let T be an $\alpha g^{\#}\psi$ -OS of (P, τ)) such that $Y \subseteq T$.

 $\Rightarrow X \subseteq T$.

As X is $\alpha^* g^{\#} \psi$ -CS and $\alpha \operatorname{cl}(X) \subseteq T$,

Then $\alpha \operatorname{cl}(Y) \subseteq \alpha \operatorname{cl}(\alpha \operatorname{cl}(X)) = \alpha \operatorname{cl}(X) \subseteq T$.

Hence, Y is $\alpha^* g^\# \psi$ -CS of (P, τ).

Remark 4.8: The above diagram shows the relationships established between

 $\alpha^* g^{\#} \psi$ -closed sets and some other sets in theorems mentioned above.

A →B represents A implies B but not conversely and

A ↔ B represents A and B are independent of each other.

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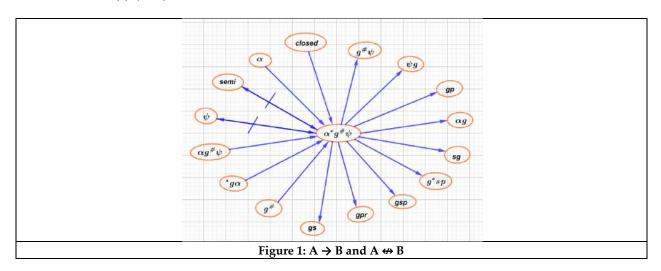
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RESEARCH ARTICLE

Further Studies on Continuity of γ *grw Closed Set with Grill

M.K. Eswarlal¹, Dr. R. Bhavani²

¹Assistant Professor, Department of Mathematics, Sourashtra College, (MKU22PFOS10531), Madurai Kamaraj University, Madurai, Tamil Nadu, India.

²Assistant Professor and Research Supervisor, PG and Research Department of Mathematics, MTN College, (Affiliated to MKU) Madurai-4.

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*Address for Correspondence

M.K. Eswarlal

Assistant Professor, Department of Mathematics, Sourashtra College, (MKU22PFOS10531), Madurai Kamaraj University, Madurai, Tamil Nadu, India. Email: eswarlalmaths@gmail.com



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ABSTRACT

This research paper deals with continuous (Cts) functions in Grill Topological space (GTS). Further investigated by introducing G γ * grwc (Grill γ *generalized regular weakly closed) set, useful results are derived and it is a Continuous study of [3] "On continuity of γ * grwc sets."

Keywords: G γ *grwc set.

INTRODUCTION

The study of shapes and spaces is known as topology and its characteristics are completely kept during steady distortion, like bending and stretching but not tearing. After the introduction of Generalized closed set by Levine (1970) it has become one of an interesting area in the field of topology. Choquet, one who explored Grill as a Tool like nets and filters. This paper deals with Continuity of G γ * grwc sets. $\mathcal{C}\iota$ and I represents closure and interior of a set. Review of literature and preliminaries, $G\gamma^*$ grwc continuous function and conclusion were presented in section 2,3 and 4 respectively.

Review of Literature and preliminaries

Related to study area a few literatures, preliminaries and definition were presented. In 1947 Choquet introduced Grill, since then many topologists started to do more research work On continuous function using GTS. Dhananjoy mandal and M.N.Mukerjee (2012) [2] have studied a new class of generalized closed sets in TS X and defined in terms of grill G on X and Presented the characterization of Regular and Normal spaces. M.O. Mustafa and





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R.B.Esmaeel(2021) [8] have conducted a research work on" some properties in Grill topological open and closed sets" and given its relationship with continuous function Gg-continuous function, Gg strongly continuous function, Gg-Irresolute function. Nagarajan Kalaivani, Khaleel Fayaz UrRahaman, Lanka cepova and Robert cep (2022) [9] have investigated GS $_{\beta}$ continuous function along with this the theory of GS $_{\beta}$ continuous mapping was introduced and derived properties in GTS. M.K. Eswarlal and R.Bhavani (2023) [3] have introduced γ *grwc set and studied its Continuity in a Topological spaces.

Basic concepts of Topology

Definition 2.1 Closure (Cι) and Interior (I) of a set.[3]

Given a subset A of a topological space X, $C\iota(A)$ is defined as the intersection of all closed sets containing A and I(A) is defined as the union of all open sets contained in A.

Definition 2.2

Let (X,τ) be TS and a subset A of X is said to be

- (i) α open [7] if $A \subseteq I(C\iota(I(A)))$ and α closed [7] set if $C\iota(I(C\iota(A))) \subseteq A$.
- (ii) Weakly closed (w-closed) [10], if $Cu(A) \subseteq H$ whenever $A \subseteq H$ and H is semi-open in (X, τ) .
- (iii) γ open [4], if $A \subseteq I(C\iota(A)) \cup C\iota(I(A))$ and γ closed [4] if $C\iota(I(A)) \cap I(C\iota(A)) \subseteq A$.
- (iv) Semi open set [5] if $A \subseteq C\iota(I(A))$ and semi closed set [5] if $I(C\iota(A)) \subseteq A$.

Definition 2.3

Let (X,τ) be TS &A \subseteq X is called

- (i) Generalized closed set (g closed) [6] if $C\iota(A) \subseteq U$ whenever $A \subseteq U$ and U is open in (X, τ) .
- (ii) Generalized α closed (g α closed)[5] if α cl(A) \subseteq H and H is α open in (X, τ)
- (iii) Regular generalized weakly set (rgw closed) [12], if $C_l(I(A)) \subseteq U$ whenever $A \subseteq U$ and U is regular semi open in X.
- (iv) sg closed [1], if $sC\iota(A) \subseteq U$ whenever $A \subseteq U$ and U is semi open in X.

Preposition 2.1

- (i) EveryClosed, semi closed, γ , α , and w are γ *grw closed.
- (ii) Every g, sg, rgw and g α closed are γ^* grw closed.

Definition 2.4

Grill:[11] A collection G of non-empty subsets of a space X is called a Grill on X if

- (i) $A \in G$ and $A \subseteq B \subseteq X$ implies $B \in G$, and
- (ii) A, $B \subseteq X$ and $A \cup B \in G$ implies $A \in G$ or $B \in G$.

Definition 2.5

[11] Let G be a Grill on a topological space (X,τ) . A mapping Φ : $P(x) \to P(x)$. Denoted by $\Phi_G(A,\tau)$ (for $A \in OP(x)$) or $\Phi_G(A)$ or simply by $\Phi(A)$ is defined by

 $\Phi_G(A) = \{x \in X : A \cap U \in G, \text{ for all } U \in \tau(x)\}.$

G γ *grw ccontinuous(Cts)function.

G γ *grwc continuous function is introduced and studied.

Definition 3.1

Let G be a Grill on X and A be a subset of X in a topological space (X,τ) , is said to be $G \gamma^* grwcif \Phi_G(A) \subseteq H$ whenever $A \subseteq H$ and H is γ closed in X.

Example 3.1

Let $X = \{j, o, v, z\}$, $\tau = \{\emptyset, X, \{j\}, \{o\}, \{j, o\}, \{j, o, v\}\}$ and

 $G = \{\emptyset, X, \{j\}, \{o\}, \{j, o\}, \{j, v\}, \{j, z\}, \{j, o, v\}, \{j, o, z\}, \{j, v, z\}\}$

Closed sets of (X, τ) are $\{\emptyset, X, \{z\}, \{v, z\}, \{o, v, z\}, \{j, v, z\}\}$

 $G\gamma^*grw$ closed sets of (X,τ) are $\{\emptyset,X,\{j\},\{o\},\{v\},\{j,v\},\{j,z\},\{o,v\},\{o,z\},\{v,z$





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 $\{j,o,z\},\{o,v,z\},\{j,v,z\}\}$

Preposition 3.1

Every closed set is $G \gamma$ *grw closed.

Proof:

Let Abe a closed set in X and A \subseteq H.

since A is closed this implies that $C\iota(A) \subseteq A \subseteq H$, where H is γ closed.

But $\Phi_G(A) \subseteq \mathcal{U}(A) \subseteq H$. (i.e) $\Phi_G(A) \subseteq H$. Thus, A is $G \gamma^*$ grw closed.

Remark 3.1

The converse of preposition 3.1 is not true.

Example 3.2

Let $X = \{j, o, v, z\}$, $\tau = \{\emptyset, X, \{j\}, \{o\}, \{j, o\}, \{j, o, v\}\}$ and

 $G = \{\emptyset, X, \{j\}, \{o\}, \{j, o\}, \{j, v\}, \{j, z\}, \{j, o, v\}, \{j, o, z\}, \{j, v, z\}\}$

Take A = $\{o\}$, Here $\{o\}$ is not closed, but it is G γ *grw closed.

Preposition 3.2

- (i) Every α , semi closed, γ and w are G γ *grw closed.
- (ii) Every rgw, g α , sg and g closed are $G\gamma^*$ grw closed.

Remark 3.2

The converse of Preposition 3.1 & 3.2 areneed be not true.

Definition 3.2

A function $f:(X,\tau,G)\to (Y,\sigma)$ is called $G\gamma^*grwc$ continuous (Cts) if the inverse image of every closed set in (Y,σ) is $G\gamma^*grw$ closed in (X,τ,G) .

Theorem 3.1

EveryCts function is $G\gamma^*grwcCts$.

Proff:

Let $f:(X,\tau,G) \rightarrow (Y,\sigma)$ be a Cts map.

Let H be a closed set in (Y , σ). Then f⁻¹ (H) is closed in (X , τ , G).

Since every closed set is $G \gamma^*$ grw closed. (i. e) $f^{-1}(H)$ is a $G \gamma^*$ grw closed.

Therefore f is Gγ*grwcCts.

Remarks 3.3

Converse of Th 3.1 need not be true.

Example 3.3

Let $X = \{j, o, v, z\}$, $\tau = \{\emptyset, X, \{j\}, \{o\}, \{j, o\}, \{j, o, v\}\}, \sigma = \{\emptyset, Y, v\}$ and

 $G = \{\emptyset, X, \{j\}, \{o\}, \{j, o\}, \{j, v\}, \{j, o, v\}, \{j, o, z\}, \{j, v, z\}\}\$

Let $f:(X, \tau, G) \rightarrow (Y, \sigma)$ be defined as f(j) = j, f(o) = o, f(v) = v and f(z) = z.

. $f^{-1}(j,o,z) = \{j,o,z\}$. Which is not closed in X. That is f is notCts.

But $\{j,o,z\}$ is $G \gamma^*$ grw closed. Therefore f is $G \gamma^*$ grwcCts.

Theorem 3.2

- (i) Every $\alpha cCtsis G\gamma^*grwcCts$.
- (ii) Every semi closedCtsis $G\gamma^*$ grwcCts.
- (iii) Every wcCtsis Gγ*grwcCts.
- (iv) Every γ cCtsis $G\gamma^*$ grwcCts.

Remarks 3.2

Proof of Th 3.2 (i) to (iii) follows from above definitions and known results and its converse need not be true.

Example 3.3

Let $X = \{j, o, v, z\}$, $\tau = \{\emptyset, X, \{j\}, \{o\}, \{j, o\}, \{j, o, v\}\}, \sigma = \{\emptyset, Y, j\}$ and

 $G = \{\emptyset, X, \{j\}, \{o\}, \{j, o\}, \{j, v\}, \{j, z\}, \{j, o, v\}, \{j, o, z\}, \{j, v, z\}\}$

Let $f:(X, \tau, G) \to (Y, \sigma)$ be defined as f(j) = o, f(o) = j, f(v) = v and f(z) = v.

- (i) $f^{-1}(o, v, z) = \{j, v\}$. Which is not α closed. That is f is not α Cts. Therefore f is $G \gamma^* grwc$ Cts.
- (ii) $f^{-1}(j, o, z) = \{j, o, v\}$ and $f^{-1}(j, o) = \{j, o\}$ Which is not w closed. That is f is w Ctsbut not $G\gamma^*g$ rwcCts.





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Let $X = \{j, o, v, z\}$, $\tau = \{\emptyset, X, \{j\}, \{o\}, \{j, o\}, \{j, o, v\}\}$, $\sigma = \{\emptyset, Y, j, v\}$ and

 $G = \{\emptyset, X, \{j\}, \{o\}, \{j, o\}, \{j, v\}, \{j, z\}, \{j, o, v\}, \{j, o, z\}, \{j, v, z\}\}$

Let $f:(X, \tau, G) \to (Y, \sigma)$ be defined as f(j) = j, f(o) = o, f(v) = v and f(z) = z.

- (i) $f^{-1}(j, o, z) = \{j, o, z\}$. Which is not semi closed .That is f is not semi closed Cts. Therefore f is $G \gamma^* grwCts$.
- (ii) $f^{-1}(j, o, z) = \{j, o, z\}$ and $f^{-1}(j, o) = \{j, o\}$ Which is not both γ as well as $G \gamma^*$ grw closed . Therefore fis not γ and $G \gamma^*$ grwcCts .

Remark 3.3

Thus fig (i) depicts the relationship between $G\gamma^*$ grwcCtswith other sets.

Theorem 3.3

- (i) Every gcCtsis $G\gamma^*$ grwCts.
- (ii) Every sgcCtsis $G\gamma^*$ grwCts.
- (iii) Every rgwcCts isGγ*grwCts.
- (iv) Every g $\alpha cCtsis G\gamma^*grwCts$.

Remarks 3.4

Proof of Th 3.3 (i) to (iv) follows from above definitions and known results and its converse need not be true.

Example 3.4

Let $X = \{j, o, v, z\}$, $\tau = \{\emptyset, X, \{j\}, \{o\}, \{j, o\}, \{j, o, v\}\}, \sigma = \{\emptyset, Y, j\}$ and

 $G = \{\emptyset, X, \{j\}, \{o\}, \{j, o\}, \{j, v\}, \{j, z\}, \{j, o, v\}, \{j, o, z\}, \{j, v, z\}\}$

Let $f:(X, \tau, G) \to (Y, \sigma)$ be defined as f(j) = 0, f(o) = j, f(v) = v and f(z) = v.

- (i) $f^{-1}(o, v, z) = \{j, v\}$. Which is not g closed. That is f is not g continuous. Therefore f is $G \gamma^* grwcCts$.
- (ii) $f^{-1}(j, o, z) = \{j, o, z\}$ and $f^{-1}(j, o) = \{j, o\}$ Which is not both sg as well as $G \gamma^* grw$ closed . (i,e) f is not sgand $G \gamma^* grw$ cCts.
- (iii) $f^{-1}(j, o, z) = \{j, o, z\}$ and $f^{-1}(j, o) = \{j, o\}$ Which is not both $g \alpha$ as well as $G \gamma^* grw$ closed. Therefore f is not $g \alpha$ and $G \gamma^* grw$ cCts.
- (iv) Let $\sigma = \{\emptyset, Y, \{j\}, \{j, z\}\}\$ and $f^{-1}(j, v, z) = \{j, v\}, f^{-1}(o, v) = \{j, v\}.$ Which is not rgw closed. That is f is not rgw continuous. That is f is $G\gamma^*grwcCts$.

Remark 3.5

fig (ii) shows the association of $G\gamma^*grwcCts$ with other sets

CONCLUSION

 $G\gamma^*$ grw Cts function was introduced and investigated its link between other sets were presented. This study can be extended to any topological space.

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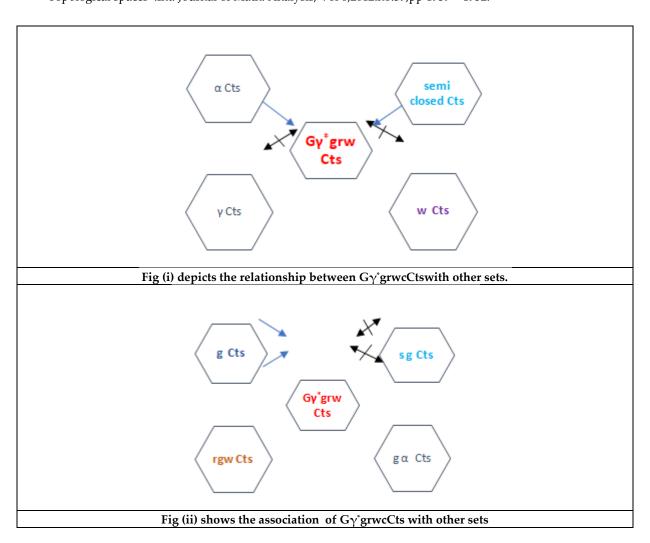
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RESEARCH ARTICLE

Cyclotide Induced Neuronal Cell Death **Prevents** Hydroxydopamine Neurotoxin

Shobana C*, Usharani B and Rohini D

Department of Biochemistry, School of Life Sciences, Vels Institute of Science, Technology and Advanced Studies, (Affiliated to Deemed University) Tamil Nadu, India.

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*Address for Correspondence Shobana C

Department of Biochemistry, School of Life Sciences, Vels Institute of Science, Technology and Advanced Studies, (Affiliated to Deemed University) Tamil Nadu, India. Email: shobana.sls@velsuniv.ac.in



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ABSTRACT

Neuronal cell death is a major factor in the initiation and progression of Parkinson's disease (PD). Reactive oxygen species (ROS) overproduction is one of the primary causes of neuronal death, hence drugs that decrease oxidative stress-dependent neuronal death as a preventive measure against Parkinson's disease (PD) may be promising. However, cyclotide's preventive effects have not been studied using both in vitro and in vivo models of Parkinson's disease. Therefore, in order to investigate the efficacy of cyclotide against 6-hydroxydopamine (6-OHDA)-dependent neuronal cell death, we employed immortalized hypothalamic neurons (GT1-7 cells). First, we found that cyclotide prevents 6-OHDA-dependent neuronal cell death by inhibiting ROS overproduction in GT1-7 cells. The cytoprotective effect of cyclotide was largely abolished by verapamil, an OCTN1 inhibitor that inhibits cyclotide absorption. According to these results, cyclotide or foods containing it may be a helpful tactic to postpone the start and progression of Parkinson's disease (PD).

Keywords: Cyclotide; neuronal cell death; 6-OHDA; oxidative stress; functional food

INTRODUCTION

Parkinson's disease (PD) is one of the most common progressive neurodegenerative illnesses. Parkinson's disease (PD) affects both the central nervous system and the body parts that are innervated by diseased nerves. This results in altered postural reflexes, stiffness in the muscles, akinesia, and resting tremor. In the initial stages of Parkinson's disease (PD), sluggish movements and trouble walking are frequently observed. People who have the disease gradually become bedridden or unable to move freely, requiring them to use a wheelchair [1,2].





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Amantadine hydrochloride, anticholinergic drugs, and the dopamine precursor L-dopa are commonly used in the treatment of Parkinson's disease (PD) patients in order to compensate for the dopamine depletion brought on by the death of dopaminergic neurons. Despite the moderate efficacy of these drugs [3], there is now no established method to stop the onset and progression of Parkinson's disease. Parkinson's disease (PD) has no known exact etiology, although elevated inflammatory response and loss of dopaminergic neurons in the substantia nigra of the midbrain are established risk factors. Lesion-related microglia are thought to be the main cause of Parkinson's disease (PD) [4,5]. Treating mice or neurons with 6-hydroxydopamine (6-OHDA) or N-methyl-4-phenylpyridinium iodide (MPP+) has been used in basic research because these substances can produce models that mimic oxidative stress-mediated dopaminergic neuronal degeneration and enhanced microglial inflammation [6,7]. Moreover, disturbance of the hypothalamic-pituitary-adrenal axis is said to have an impact on the development and progression of Parkinson's disease (PD) and govern the synthesis of numerous hormones [8]. The comparison of the plasma concentrations of adrenocorticotropic hormone and nocturnal growth hormone in Parkinson's disease (PD) patients with those of healthy controls indicates possible abnormalities in the hypothalamus [9].

A reduction in the quantity of neurons in the hypothalamus of Parkinson's disease patients was found in one study [10]. Furthermore, a strong positive correlation has been observed between the clinical stage of Parkinson's disease and a decrease in orexin-containing neurons, a hormone that concentrates melanin in the hypothalamus [11]. These preliminary findings suggest that inhibiting neuronal death in the hypothalamus may prevent the onset and progression of Parkinson's disease. One of these potentially beneficial herbs is Clitoria ternatea, a key ingredient in the brain tonic medhya rasayan, which is used to treat neurological illnesses. Ayurvedic medicine has been used traditionally in India from ancient times. This study supports Indian medicine by highlighting the plant's significance as a brain remedy. Clitoria ternatea has a potent brain-stimulating effect, unlike other plants [12]. Moreover, C. ternatea has long been used in traditional medicine, particularly as a supplement to improve cognitive function and lessen the symptoms of numerous illnesses like fever, inflammation, discomfort, and diabetes [13]. This study assessed the antiparkinson's activity of cyclotide, the active ingredient in Clitoria ternatea. Previous work on the analysis of cyclotide insilico and invitro served as the foundation for the current investigation. Despite the fact that cyclotide has been linked to a number of biological processes, its ability to prevent 6-OHDA-dependent hypothalamic neuronal cell death has not been studied. Therefore, we used immortalized hypothalamic neurons and GT1-7 cells in this investigation to investigate the effectiveness of cyclotide on 6-OHDA-dependent neuronal cell death. Additionally, we ascertained cyclotide's impact on the generation of ROS in response to 6-OHDA as well as its effectiveness in preventing neuronal cell death caused by 6-OHDA.

MATERIALS AND METHODS

Chemicals and Reagents

MPP+ was supplied by Cayman Chemical. Promega Corporation was the CellTiter-Glo® 2.0 provider. Fujifilm Wako Pure Chemical Corporation in Tokyo, Japan was the seller of DMEM/Ham's Nutrient Mixture F-12 (Dulbecco's Modified Eagle's Medium/Ham's Nutrient Mixture). The 2',7'-dichlorodihydrofluorescein diacetate (H2DCFDA) was supplied by Merck KGaA. The FastGeneTM RNA Basic kit was provided by Nippon Genetics Co., Ltd.; the PrimeScriptTM RT master mix (Perfect Real Time) was provided by Takara Bio; and the THUNDERBIRD® Next SYBR® qPCR mix was provided by Toyobo.

Culture of Cells

For the investigation, immortalized hypothalamic neurons, or GT1-7 cells, were employed. 10% fetal bovine serum was added to DMEM/Ham's-F12, the growth media used for GT1-7 cells. After being treated with trypsin (Fujifilm Wako Pure Chemicals), the cells were suspended in a serum-free medium, plated, and cultured at 37°C in a humidified incubator with 7% CO2 [15].





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Determination of cell viability.

Cell viability was evaluated, same as it was in our previous studies [20, 21]. To put it briefly, $3.0 \times 104 \text{ GT1-7}$ cells were seeded per well onto 96-well growth plates using 200 µL of culture medium. After a 24-hour preincubation period, the medium was supplied with either 6-OHDA (final concentrations ranging from 0 µmol/L to 80 µmol/L) or MPP+ (final concentrations ranging from 0 mmol/L to 8 mmol/L). As an alternative, the cells were preincubated for 24 hours before being treated with cyclotide (final concentrations ranging from 0 mmol/L to 1.0 mmol/L) for 10 min. The medium was subsequently supplemented with either MPP+ (final concentration: 4 mmol/L) or 6-OHDA (final concentration: 40 mmol/L). The cells' vitality was evaluated using the luminous reagent CellTiter-Glo® 2.0, which quantifies intracellular ATP, following a 24-hour incubation period. Assuming that cyclotide is ingested as a daily meal or supplement, it was pretreated to GT1-7 cells in this investigation. Its efficacy was evaluated. Furthermore, despite discrepancies in reporting, the blood concentration of cyclotide in humans is roughly 0.2 mmol/L, which is comparable to the concentration of cyclotide used in this experiment [22].

ROS Level Measurement

GT1: Seven cells were cultured in black 96-well microplates with 3.0 \times 104 cells per well for a whole day. Next, the cells were exposed to the ROS indicator H2DCFDA (10 μ mol/L) for 60 minutes. Subsequently, cyclotide was added to the media (final concentration: 40 μ mol/L) and the cells were treated with it (final concentrations ranging from 0 mmol/L to 1.0 mmol/L). After an hour, the ROS levels were measured using a microplate reader (excitation: 480 nm, emission: 530 nm).

Instantaneous Analysis of Reverse Transcription Polymerase Chain Reaction (RT-PCR)

Total RNA was isolated from GT1-7 cells according to the manufacturer's instructions using the FastGeneTM RNA Basic kit. Samples were reverse-transcribed using the PrimeScript RT master mix. The resulting cDNA was analyzed in real-time PCR experiments with THUNDERBIRD Next SYBR qPCR mix using CFX ManagerTM software (Version 3.1) on a Bio-Rad CFX96TM realtime system (Hercules, CA, USA). The specificity was verified by electrophoretic analysis of the reaction products with template- or reverse transcriptase-free controls. Glyceraldehyde-3-phosphate dehydrogenase (Gapdh) cDNA was employed as an internal standard to adjust the total RNA concentration in every experiment. Primers were designed using the Primer-BLAST website (https://www.ncbi.nlm.nih.gov/tools/primer-blast/).Primer sequences are shown in Supplementary Figure S1.

Statistical Analysis

All data are expressed as mean \pm using the standard error of the mean (S.E.M.). One-way analysis of variance (ANOVA) was used to examine differences between three or more groups, or between two groups. For unpaired data, Dunnett's test or Student's t-test were used, respectively. A difference was considered significant if it was p < 0.05 (* or # p < 0.05, ** or ## p < 0.01). Details on the symbols are provided by the figure legends. The number of samples (n) is also indicated in each image.

RESULTS

Efficacy of Cyclotide on 6-OHDA-Induced Neuronal Cell Death

The development and course of Parkinson's disease are significantly influenced by neuronal cell death [4,5]. In addition to producing a variety of Parkinson's disease-like symptoms when injected into the brains of test animals, 6-OHDA also acts as a neurotoxic in cellular experimental setups, leading to the death of neurons [6,23]. We started by evaluating the survival of GT1-7 cells, an immortalized line of mouse hypothalamus neurons, that were only exposed to 6-OHDA. Cell viability decreased in a concentration-dependent manner in response to treatment of 20, 40, 60, and 80 μ mol/L concentrations of 6-OHDA (Figure 1A). The groups' respective post-treatment viabilities were 73.7 \pm 2.7, 47.4 \pm 2.7, 33.2 \pm 2.3, and 23.9 \pm 1.6% (mean \pm S.E.M., n = 4). All groups shown a noteworthy decrease in comparison to the ultrapure water-treated control group. Next, we investigated the effect of cyclotide pretreatment using a 40 μ mol/L final concentration of 6-OHDA on the 6-OHDA-induced decrease in GT1-7 cell viability. Pretreatment with





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0.5 mmol/L and 1.0 mmol/L of cyclotide restored cell viability to $77.1 \pm 2.5\%$ and $82.8 \pm 0.8\%$ (mean \pm S.E.M., n = 4) respectively, effectively mitigating the 6-OHDA-dependent decrease in cell viability. The protective effect was concentration-dependent. Under these circumstances, cyclotide therapy alone had little effect on cell viability (Figure 1C). Likewise, MPP+ functions as a neurotoxic in experimental settings that are both in vivo and in vitro [7]. We looked into if cyclotide also prevented the death of neurons triggered by MPP+. Reduced cell viability was seen in GT1-7 cells treated with 1.0-8.0 mmol/L MPP+ (Figure 1D). The MPP+-dependent loss in cell viability was considerably mitigated by cyclotide; after pretreatment with 0.5 mmol/L and 1.0 mmol/L of cyclotide, respectively, cell viability reached $53.0 \pm 1.3\%$ and $55.9 \pm 0.5\%$ (mean \pm S.E.M., n = 4). (Figure 1E). Numerous investigations suggest that the endoplasmic reticulum (ER) stress response plays a crucial role in the neuronal cell death caused by 6-OHDA [24, 25]. Our lab has previously shown that enhanced ER stress-related gene expression in GT1-7 cells produced by 6-OHDA may be suppressed by the antioxidant peptide carnosine and an antioxidant proteinthioredoxin-albumin fusion [26,27]. Treatment with 40 µmol/L of 6-OHDA in this study elevated the expression of genes linked to ISR, especially the significant elevation of growth-arrest and DNA-damage-inducible gene 34 (Gadd34) and CCAAT-enhancer-binding protein homologous protein (Chop). In particular, compared to the control group, Chop and Gadd34 mRNA increased by 6.72 ± 0.08-fold and 4.60 ± 0.22-fold (mean ± S.E.M., n = 3), respectively. In addition, 6-OHDA treatment significantly increased the levels of ER degradation-enhancing α mannosidase (Edem), binding immunoglobulin protein (Bip), activating transcription factor 4 (Atf4), inositolrequiring transmembrane kinase/endoribonuclease 1α (Ire1α), protein disulfide isomerase (Pdi), and glucoseregulated protein 94 (Grp94). On the other hand, the overexpression of these genes was inhibited by cyclotide pretreatment, especially Chop (to 2.50 ± 0.15 -fold) and Gadd34 (to 2.14 ± 0.04 -fold) mRNA (mean \pm S.E.M., n = 3). According to these results, cyclotide suppresses the elevated expression of genes linked to ER stress, hence preventing 6-OHDA-induced neuronal cell death (Figure 2).

Efficacy of Cyclotide on 6-OHDA-Induced Reactive Oxygen Species (ROS) Production

In recent studies, it was found that the antioxidant peptide carnosine and the antioxidant protein-thioredoxinalbumin fusion both successfully reduced the increased expression of genes linked to ER stress that 6-OHDA caused in GT1-7 cells [26, 27]. In the present investigation, 40 μ mol/L of 6-OHDA induced an upregulation of ISR-related gene expression, most notably CCAAT-enhancer-binding protein homologous protein (Chop) and growth-arrest and DNA-damage-inducible gene 34 (Gadd34). In particular, compared to the control group, Chop and Gadd34 mRNA levels increased by 6.72 \pm 0.08-fold and 4.60 \pm 0.22-fold (mean \pm S.E.M., n = 3), respectively. Furthermore, after 6-OHDA treatment, there were notable increases in the following processes: binding immunoglobulin protein (Bip), ER degradation enhancing α mannosidase (Edem), activating transcription factor 4 (Atf4), inositol-requiring transmembrane kinase/endoribonuclease 1α (Ire 1α), protein disulfide isomerase (Pdi), and glucose-regulated protein 94 (Grp94). On the other hand, cyclotide pretreatment inhibited the overexpression of these genes, specifically reducing the mRNA levels of Gadd34 (down to 2.13 \pm 0.04-fold) and Chop (down to 2.52 \pm 0.15-fold) (mean \pm S.E.M., n = 3). These results clearly suggest that cyclotide has the ability to inhibit the overexpression of genes linked to ER stress that is brought on by 6-OHDA.

Involvement of OCTN1 in Cytoprotective Effects of Cyclotide

The protein known as OCTN1, a membrane transporter, is in charge of identifying and moving endogenous or exogenous substances into and out of cells. OCTN1 is essential for cyclotide absorption, as demonstrated by a number of investigations on mice and cultured cells [15,16]. Therefore, we looked into whether verapamil, a well-known OCTN1 inhibitor, would neutralize the protective effect of cyclotide on cells [30]. As expected, cyclotide successfully counteracted the cell viability loss caused by 6-OHDA (Figure 4). However, the protective effect of cyclotide was greatly diminished when cells were pretreated with verapamil. Notably, verapamil administration by itself did not significantly affect ROS generation or cell viability in this experimental setup (Supplementary Figure S2). These results strongly imply that the cytoprotective action of cyclotide in GT1-7 cells is dependent on its entry into the cell through OCTN1.





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DISCUSSION

In this study, we examined the protective effect of cyclotide against 6-OHDA-induced neuronal cell death using GT1-7 cells as immortalized hypothalamic neurons. Our results show that cyclotide significantly reduces the overexpression of ER stress-related proteins, including Chop and Gadd34, in GT1-7 cells as well as 6-OHDA-induced cell death. The pathophysiology and exacerbation of a number of diseases, including cancer, pneumonia, and neurological disorders, are significantly influenced by the ER stress response, which is mediated by protein kinase R-like ER kinase (PERK), inositol-requiring enzyme-1 (IRE1), and ATF6 [33, 34]. When these sensors are activated, the cell begins to undergo signaling processes, such as PERK phosphorylating eIF2's α subunit, which affects ATF4 translation in turn.ATF4 in turn initiates the transcription of GADD34 and CHOP, which are involved in the induction of cell death [33, 34]. We suggest that cyclotide prevents 6-OHDA-dependent neuronal cell death by inhibiting the expression of proteins linked to ER stress and cell death, specifically Gadd34 and Chop, as well as the transcription factor that is upstream of these factors, Atf4. In conclusion, our groundbreaking discovery emphasizes how cyclotide suppresses the ER stress response, hence inhibiting 6-OHDA-induced neuronal cell death and suggesting the drug's potential as a PD preventative. Since 6-OHDA is known to cause apoptosis in neuronal cells [26, 29], we will be investigating a more thorough way of triggering cell death in subsequent studies.

In our work, the OCTN1 inhibitor verapamil inhibited the cytoprotective effect of cyclotide; nevertheless, it is important to highlight that this protective effect was not completely eliminated by verapamil pretreatment. Furthermore, verapamil by itself demonstrated cytotoxicity at values higher than 100 µmol/L (Supplementary Figure S2), which raises questions about its suitability for use in research. As a result, future research should identify additional pathways that contribute to cyclotide's cytoprotective action in addition to OCTN1. One of the main causes of neuronal cell death and exaggerated inflammatory responses in Parkinson's disease (PD) is oxidative stress, which is marked by elevated generation of reactive oxygen species [28,29,35]. Clinical research has shown a direct link between the generation of ROS and Parkinson's disease. Cyclotide appears to be a promising treatment option for Parkinson's disease (PD) since it inhibits the creation of excessive ROS or oxidative stress. This idea is in line with earlier research that shown the effectiveness of antioxidants, including N-acetylcysteine and coenzyme Q10, in reducing Parkinson's disease symptoms in both human and animal trials. Moreover, cyclotide has proven effective in stroke and Alzheimer's disease animal models, exhibiting neuroprotective and antioxidant properties [17, 18]. Its neuroprotective qualities have also been shown in cultured cells, where it prevents amyloid-beta and anticancer medications from injuring neuronal cells. While cyclotide's effectiveness in treating a variety of neurological conditions has been studied, there have been no reports of studies examining its effectiveness in PD animal models or 6-OHDA-induced neuronal cell death, with the exception of one clinical study that found PD patients' blood levels of cyclotide were lower than those of controls. These results highlight the potential importance of adding cyclotides to meals or supplements as a critical prophylactic against the onset and progression of Parkinson's disease.

Through its antioxidant qualities, cyclotide, we found in our inquiry, protects against 6-OHDA-dependent neuronal cell death. Based on these findings, we propose that cyclotide may be a viable treatment option for delaying the onset and progression of Parkinson's disease (PD). In the future, we want to investigate more thoroughly how well cyclotide works in animal models of Parkinson's disease. GT1-7 cells were treated with the indicated concentrations of 6-hydroxydopamine (6-OHDA) (µmol/L) or N-methyl-4-phenylpyridinium (MPP+) (mmol/L) and incubated for 24 h (A,D). GT1-7 cells were pretreated with cyclotide (0.06–1.0 mmol/L) and then incubated in the absence (Control) or presence of 6-OHDA (40 µmol/L) mmol/L)alone and cultured for a further 24 h (C). Cell viability was measured using CellTiter-Glo® 2.0. Values represent mean \pm S.E.M. (n = 4). ** p < 0.01, vs. Control; ## p < 0.01, vs. 6-OHDA (40 µmol/L) alone or MPP+ (4 mmol/L) alone. GT1-7 cells were pretreated with cyclotide (1.0 mmol/L) and then incubated in the absence (Control) or presence of 6-OHDA (40 µmol/L) for 24 h. After total RNA extraction from GT1-7 cells, cDNA was synthesized, and real-time RT-PCR was performed using primer pairs that specifically amplify Chop, Gadd34, Atf4, Bip, Ire1 α , Pdi, Edem, and Grp94. Values were normalized to Gapdh and expressed relative to control. Values represent mean \pm





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S.E.M. (n = 3). * p < 0.05, vs.Control; ** p < 0.01, vs. Control; # p < 0.05, vs. 6-OHDA (40 μ mol/L) alone; ## p < 0.01, vs. 6-OHDA(40 μ mol/L) alone. GT1-7 cells were pretreated with a reactive oxygen species (ROS) indicator, 2',7'-dichlorodihydrofluorescein diacetate (H2DCFDA)(10 μ mol/L) for 60 min (A,B). Cells were then treated with 6-OHDA (20–80 μ mol/L) and culturedfor 1 h (A). GT1-7 cells were pretreated with cyclotide (0.06–1.0 mmol/L) and then incubated in the absence (Control) or presence of 6-OHDA (40 μ mol/L) for 1 h (B). GT1-7 cells were treated with cyclotide (0.06–1.0 mmol/L) alone and cultured for 1 h (C). ROS levels were measured using fluorescence microplate reader. Values represent mean •} S.E.M. (n = 4). ** p < 0.01, vs. Control;## p < 0.01, vs. 6-OHDA (40 μ mol/L) alone. GT1-7 cells were pretreated with verapamil (100 μ mol/L) for 60 min. After replacement with fresh medium, cells werethen pretreated with cyclotide (0.5 or 1.0 mmol/L) and incubated in the absence (Control) or presence of 6-OHDA (40 μ mol/L) for 24 h. Cell viability was measured using CellTiter-Glo® 2.0. Valuesrepresent mean ± S.E.M. (n = 4). Not Significant (n.s.), ** p < 0.01, Control vs. verapamil (100 μ mol/L).

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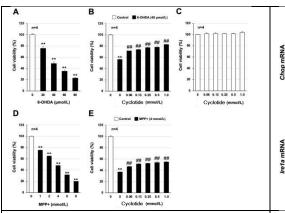
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Cyclotide (mmolif)

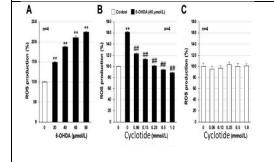
(Relative expression)

Cyclotide (mmolif)

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Figure 1. Neuroprotective effect of cyclotide

Figure 2. Cyclotide suppresses the 6-hydroxydopaminedependent endoplasmic reticulum stress response



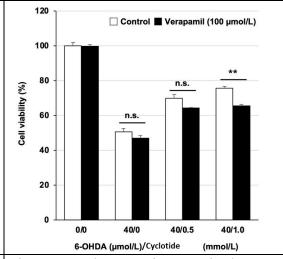


Figure 3. Antioxidant effect of cyclotide in GT1-7 cells

Figure 4. Involvement of OCTN1 in the cytoprotective effect of cyclotide.





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RESEARCH ARTICLE

A Systematic Analysis of Ethnomedicinal, Elemental, Phytochemical and Pharmacological Properties of Indian Mallow-Abutilon indicum (linn.) sweet (Malvaceae)

Rishidha VM^{1*}, Vasantharaja D² and Muthuviveganandavel V¹

Department of Zoology, Kanchi Mamunivar Government Institute for Post Graduate Studies and Research (Autonomous) (Affiliated to Pondicherry University) Puducherry, India.

²Department of Zoology, Tagore Government Arts and ScienceCollege,(Affiliated to Pondicherry University)Puducherry, India.

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*Address for Correspondence Rishidha VM

Department of Zoology,

Kanchi Mamunivar Government Institute for Post Graduate Studies and Research (Autonomous) (Affiliated to Pondicherry University)Puducherry, India.

Email: rishidharanjith94@gmail.com



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ABSTRACT

Abutilon indicum (Linn.) Sweet, often known as Indian mallow or Country mallow, is a member of the Malvaceae family of plants. This plant is indigenous to the subtropical and tropical areas of the Americas, India, Malaysia, Bangladesh, Srilanka, Pakistan and the Caribbean. There is a wide variety of uses for the plant in both conventional and alternative medicine, including the complete plant and its many components such leaves, fruits, flowers, seeds, stems, and roots. The pharmacological properties of Abutilon indicum have been documented include antibacterial, anti-inflammatory, antioxidant, hepatoprotective, nephroprotective, wound healing, antivenom, anticancer, antiarthritic, anti-Alzheimer's, hypoglycaemic, antiulcer, aphrodisiac, anti-diarrhoeal, anticonvulsant, diuretic activity, antimycotic, larvicidal, antidiabetic, analgesic and sedative activities. The Abutilon indicum milk is used to treat urinary discharges, while the roots of the plant are used to treat gout, polyuria, and hemorrhagic illnesses in Ayurveda. The Siddha medical tradition prescribes the plant for a variety of conditions, including ulcers, jaundice, leprosy, and piles. The herb is used to treat bronchitis, piles, and chest pain in Unani medicine. The various plant sections have been observed to have a wide variety of phytochemical components. This article provides an in-depth assessment of the scientific literature pertaining to the ethnomedicinal applications, elemental analysis, phytochemistry, and pharmacological characteristics of the plant *Abutilon indicum*.

Keywords: Abutilon indicum, Country mallow, Traditional medicine, Pharmacology, Phytochemistry.





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INTRODUCTION

The plant *Abutilon indicum*, known as "Thuthi" in Tamil and "Kanghi" in Hindi, is native to South Asia. Nature is a dependable and valued ally in the pharmaceutical sector. Natural treatments offer comparable advantages to synthetic ones, but without adverse side effects. The perennial plant *A. indicum*, commonly referred to as "Country Mallow," can grow as tall as 3 meters [1]. Medicinal herbs provided by nature are a gift intended to promote longevity and good health by preventing sickness. Preserving our health is one of its essential tasks. India is renowned for its extensive utilization of plants for medicinal purposes, making it one of the most diverse countries in the world in this regard. Ayurveda, Unani, and Siddha are the three main alternative medicinal practices according to Sikorska and Matlawska [2]. Throughout history, individuals in India have depended on a variety of plant elements for healing purposes. *A. indicum* exemplifies such a plant. Abutilon belongs to the Malvaceae family, with more than one fifty species of biennial herbs and occasionally trees and bushes. These plants are distributed over the tropics and subtropics of America, Africa, Asia, and Australia. Recently, scientists have become more interested in studying the species due to the fact that some plants in the genus are highly respected Ayurvedic treatments [3, 4].

Description of A. indicum

The Malvaceae family, which contains *A. indicum*, is widely distributed across India, Bangladesh, and Sri Lanka (Fig. 1). The plant is an evergreen shrub that can reach a height of three meters and features large, spherical leaves. The stem is circular with a slight purple hue. The plant has elongated petioles, pointed linear stipules, and ovate or orbicular leaves measuring 2-2.5 cm. Flowers are individual, with orange-yellow or yellow blooms on branching peduncles. Corollas are vividly yellow and blossom throughout the evening, but the calyx features oval lobes with pointed tips. Long filaments protrude from a hairy staminal tube. The color ranges from black to dark brown, with three to five seeds that are either reniform, tubercled, or minutely stellate[5].

Habitat

Found in the wild at elevations of up to 1200 meters in the sub-Himalayan region and adjacent mountain ranges [3]. **Geographical range**

India and Ceylon, encompassing tropical and subtropical regions.

Taxonomic position of A. indicum

Kingdom Plantae; **Subkingdom**: Tracheobionta; **Division**: Magnoliophyta **Class**: Magnoliopsida; **Subclass**: Dilleniidae; **Order**: Malvales; **Family**: Malvaceae; **Genus**: Abutilon; **Species**: indicum [5].

Vernacular names of A. indicum

Tamil: Tutti, Perum Tutti, Nallatutti, Paniara Hutti; English: Country Mallow, Indian mallow, Flowering maples; Malayalam: Katturam, Dabi, Uram, Tutti, Vellula Telugu: Adavibenda, Tutturu benda, Peddabenda, Tutti , Duvvenakaya, Duvvena Kayalu Kanada: Tutti Hindi: Kanghi, Kakahi, Jhampi Marathi: Akakai, Mudrika, Mudra, Petari, Karandi Odia: Pedipedika Bengali: Petari, Jhampi, Badela Guajarati: Khapat, Kansi, Kamsaki, Dabali Panjabi: Kangi, Kangibooti Sanskrit: Atibala, Kotibala, Kankatika, Bhuribala, Balika Farsi: Darakhtashaan Urudu: Kanghi Arabian: Masthul Gola [6].

USES OF A. INDICUM

Traditional uses

A. indicum has been used traditionally to treat a wide range of medical conditions, and nearly every portion of the plant has been shown to have some therapeutic use. The plant's roots are used as a demulcent, diuretic, and for treating chest infections and urethritis. The mixture made from the root is used to treat fevers, strangury, haematuria, and even leprosy. It has been discovered that the leaves may be used as a fomentation on sore muscles and joints, and that they are helpful for ulcers. Toothache, sore gums, and bladder irritation can all be treated internally using a decoction made from the leaves. Several medicinal properties, including those of a diuretic,





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astringent, a fever reducer, and an anthelmintic, are attributed to the bark. The seeds are employed as a treatment for piles, a laxative, an expectorant, and a remedy for cystitis, gleets, and gonorrhoea [7-10].

Ethnomedical uses

Numerous studies have looked into the ethnomedical usage of *A. indicum* and identified a wide variety of applications for it. According to ethnomedical research, A. indicum leaves have been used to cure a wide range of conditions, including piles, leg discomfort, eye and mouth infections [11], inflammation of the bladder [12], catarrhal bilious, diarrhoea [13], bronchitis, gonorrhoea [14, 15]. Flowers have been used for centuries as a variety of medicinal remedies, including as an antipyretic, a fertility booster, a cure for piles, and an antidote for gonorrhoea [16]. *A. indicum* fruit was used to treat piles, gonorrhoea, and a cough [17, 18]. The *A. indicum* seed has been used as a powerful aphrodisiac and treatment for bronchitis, piles, chronic cystitis, gleets, and gonorrhoea in traditional Indian medicine [19, 20]. Demulcent, aphrodisiac, laxative, astringent and diuretic, expectorant, anti-inflammatory, and analgesic properties have been ascribed to the stem and bark of A. indicum [14]. Treatment of bronchitis, gout, polyuria, uterine hemorrhagic discharge [15], urinary discharge, and urethritis can all be attained by using the roots [21]. Furthermore, the entire plant was utilised to cure urinary issues [22].

REPORTED ELEMENTS IN A. INDICUM

The traditional medicine derived from plants is very well known and has a rich history of apply in a wide variety of medical settings by utilising a variety of approaches to the healing of plants [23]. Since the time of Nagarjuna, a unique history has been produced in the sphere of the system of Indian medicine as a result of the utilisation of metals preparations as an essential component of the therapeutic process [24]. Ash material derived from plants may be employed in a variety of medicinal applications, and several effective procedures have been developed to regulate the quality of this material [25]. Minerals are another class of essential nutrients found in the human body. They are required not only to control the operations of the body but also to provide the body with its structure. Minerals, like vitamins, assist to control numerous processes that occur repeatedly in the body, which is why they are highly vital to our day-to-day lives. Vitamins also help regulate these activities [26]. According to the findings of Amit et al. [27], numerous types of elements may be found in the various components of A. indicum ash (Table. 1). His research uncovered information on the nine elements of sodium, potassium, calcium, magnesium, manganese, copper, and nickel [27]. In Table-1, the amount of elemental concentration that could be found in each sample was detailed. According to the findings of his research, the roost of A. indicum contains a significant quantity of salt (Na). In contrast to another parts of the plant, the leaves of the A. indicum plant have a high concentration of the minerals calcium (Ca), iron (Fe), and cobalt (Co). Despite the fact that magnesium (Mg) is more often present in stems. The seeds of A. indicum have a high concentration of the elements copper (Cu), manganese (Mn), and nickel (Ni). The best supply of potassium is found in the plant's flowers (K). In addition, a significant concentration of Na was discovered in the roots of A. indicum. There is an appropriate presence of a wide variety of minerals, each of which is critical to the production of energy within the body.

PHYTOCHEMISTRY

Many researchers have studied the phytochemical features of A. indicum and discovered its various chemical compositions.

Whole plant and Aerial Part

The whole plant, including the roots, contains abundant mucilaginous chemicals and asparagines [1, 2, 28, 65, 66]. The main chemical classes are listed in table 2. The plant's aerial section contains -sitosterol, an alkanol fraction, and an n-alkane combination. The blooms and branches of the plant may include saponins, flavonoids, and alkaloids [29, 30, 63, 64].





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Root

The roots were utilized to extract a non-drying oil rich in various fatty acids such as linoleic, oleic, stearic, palmitic, lauric, myristic, caprylic, and capric acids [2, 31, 67-69]. Additionally, the oil contained an unusual fatty acid with a C17 carbon structure, sitosterol, and amyrin obtained from unsaponifiable material (Table 3).

Leaves

The plant's leaves include steroids, sapogenins, sugars, and flavonoids. Eudesmic acid, ferulic acid, and caffeic acid were isolated from the methanolic extract of A. indicum leaves [2,3, 32, 62]. The compounds were identified using IR spectroscopy, 1H and 13C NMR, and chemical methods [61, 70-73]. Twenty one bioactive components were identified in the ethanolic leaf extract of A. indicum by GC-MS (Table 4).

Flower and Fruits

A total of seven unique flavonoid constituents were extracted and characterized from the flowers of A. indicum [33-36] (Table 5). Alantolatone and isoalantolactone, both sesquiterpene lactones, have been reported for the first time. Fruits have had both flavonoids and alkaloids in their compositions [37, 38] (Table 6).

Seeds

The seeds of the plant A. indicum produce a water-soluble galactomannan with a 2:3 molar ratio of galactose to mannose. Cis-12,13-epoxyoleic (vernolic) acid, 9,10-methylene octadec-9-enoic (sterculic) acid, and 8,9-smethylene-heptadec-8-enoic (malvalic) acid are all derived from the seed oil of the plant [39, 70]. Analyzing seed oil using TLC and GLC revealed a substantial quantity of unsaturated acids (Table 7). The primary components derived from the saturated acids were stearic acid and palmitic acid. Raffinose was discovered as the predominant sugar component in seed. Seed proteins consist of various amino acids, which collectively make up 31% of the total protein content [40].

PHARMACOLOGICAL ACTIVITIES OF A. INDICUM (Table. 8)

Antibacterial activity

With six different species of bacteria - B. subtilis, S. aureus, K. pneumoniae, P. aeuroginosa, E. coli, and Salmonella typhi—the antibacterial activity of the aqueous, ethanol, and chloroform extracts of A. indicum leaves was assessed using the disc diffusion method. An antibacterial effect was experimental in all three extracts. The ethanol extract showed the strongest antibacterial activity. The comparison study revealed that the leaves have significant antibacterial activity, surprising the researcher [41].

Anti-inflammatory effect

In the process of extraction, the whole A. indicum plant was utilized together with ethanol. An extract containing ethanol was administered to healthy Wistar rats at dosages of 250, 500, and 750 milligram per kilogram to be used as a model for examination. The ethanolic extract shown significant anti-inflammatory effects throughout the later stages (3 hours) of the disease's development [42].

Hepatoprotective function

A. indicum leaf aqueous extract was evaluated in rats for its hepatoprotective properties against hepatotoxicity produced by carbon tetrachloride and paracetamol [43]. The leaf's extract was administered at doses of 100 and 200 milligram per kilograms. Conventional analyses were conducted on many biochemical indicators, such as serum glutamic pyruvate transaminase, total bilirubin, direct bilirubin, and liver glutathione. Administering leaf extract to rats before they were exposed to carbon tetrachloride and paracetamol significantly reduced the toxicity induced by both substances. The hepatoprotective potential of *A. indicum* extract is high.

Nephroprotective effect

The term acute renal injury is increasingly more frequently used to describe acute renal failure [44]. Acute renal damage is a reversible condition marked by elevated creatinine (CR) and urea nitrogen in the blood serum. An





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ethanolic extract of *A. indicum* roots was examined for its nephroprotective properties in Wistar albino rats with gentamicin-induced acute renal failure. The *A. indicum* extract was given at doses of 150 and 300 milligram/kilogram of body weight during the experiment. The extracts therapy caused diminish in the amount of serum bilirubin and CR, as well as an increase in urine production, serum total proteins, and albumin. Histopathology was shown to be correlated with the biochemical changes. The most effective dosage of extract 16 for nephroprotection was 300 mg/kg.

Activity related to the healing of wounds

Albino rats were used to test the impact of the herb *A. indicum* on wound healing with a range of wounds, like excision, incision, and dead space wounds. Percentage of wound contraction, epithelization time in days, granuloma weight, and skin breaking strength were observed. Silver sulfadiazine was used as a reference standard. An ethanolic extract was given at a dose of 400 milligram/kilogram. The ethanol extract demonstrates significant wound healing effects by promoting early wound epithelization and increasing wound tensile strength [45].

Anti-venom efficacy

The study aimed to assess if the methanolic and hexane extracts from A. indicum leaves may work as an antivenom against Echis carinatus venom, commonly known as the saw-scaled viper. Snake venom was shown to include deleterious enzymes that restrict the efficacy of the extract. The enzymes mentioned include protease, phosphomonoesterase, phosphodiesterase, acetylcholine esterase, phospholipase A2, and hyaluronidase. The discovery of L-amino acid oxidase was also made. The methanolic extract from A. indicum leaves shown powerful antivenom properties by effectively inhibiting the majority of toxic enzymes generated by Echis carinatus [46].

Anticancer properties

Gold nanoparticles (GNPs) possess unique features such as small size, excellent biocompatibility, surface chemistry, low toxicity, and suitability for surface modification. Due to these characteristics, GNPs are a suitable option for biological applications. An alternative green approach was utilized to synthesize gold nanoparticles (AIGNPS) from A. indicum leaf extract (AILE). The experiment utilized HT-29 colon cancer cells to assess the cytotoxic effects of gold nanoparticles. The spherical A. indicum gold nanoparticles ranged in size from 1 to 20 nm, as determined by the invitro free radical scavenging research findings. The cytotoxic study strongly supports the use of environmentally friendly synthetic gold nanoparticles in advancing the treatment of colon cancer [47].

Anti-arthritic efficacy

An aqueous extract of the plant A. indicum was treated in vitro to determine its effectiveness in treating arthritis using several pharmacological models, such as protein denaturation inhibition, membrane stability enhancement, and proteinase inhibition. The effects of administering the extract at concentrations of 100 and 200 mg/mL were contrasted with those of acetyl salicylic acid at a concentration of 250 milligram per milliliter. The herbal extract of A. indicum demonstrated a significant and dose-dependent antiarthritic activity [48].

Anti-Alzheimer's properties

Alzheimer's disease is characterized by memory loss and cognitive issues, along with an increase in acetylcholinesterase levels leading to a decrease in acetylcholine levels. Beta sitosterol in A. indicum is accountable for the plant's anti-inflammatory and antioxidant properties [49]. The spectrophotometric analysis of the methanolic extracts revealed a substantial level of acetylcholinesterase inhibitory activity. Sprague-Dawley rats were administered aluminum chloride to induce Alzheimer's disease, and the methanolic extract of the complete Abutilon indicum plant was tested for its ability to inhibit memory impairment and enhance cognitive performance. The study utilized the Radial Arm Maze (RAM) and the Elevated plus Maze (EPM) to investigate animals' working memory. The rat memory evaluator was utilized to evaluate the long-term memory quality (RME). The Ellman approach was used to evaluate the level of acetylcholinesterase enzyme activity in the brain. The extract was given at doses of 400 and 600 milligram/kilogram. According to biochemical examination, the rats treated with the drug had significantly reduced levels of ACE enzyme. MEAI successfully enhanced remembrance by repairing the cognitive impairment





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caused by aluminum chloride. It may be concluded that Abutilon indicum extract has a protective effect against Alzheimer's disease induced by AlCl₃.

Hypoglycemic effect

The potential of A. indicum leaf extracts to reduce blood sugar levels in Wistar rats was investigated using aqueous solutions, petroleum ether, chloroform, and ethanol. The oxidase/peroxidase technique was used to determine the glucose concentration in the blood. The patients were administered a dosage of 400 mg/kg of the extracts [50]. Tolbutamide was used as a reference point. No significant hypoglycemic effect was seen in the pet-ether or chloroform extracts. Both aqueous and ethanol extracts shown considerable hypoglycemic effects in common rats four hours after injection, with reductions of 26.95% and 23%, respectively.

Antiulcer efficacy

The study investigated the effectiveness of the leaves of A. indicum in treating ulcers in rats produced by pylorus ligation and ethanol. Various parameters such as pH, stomach volume, total acidity, free acidity, and ulcer index were examined in the rat model induced by pylorus ligation. The ulcer index and the extent of ulceration inhibition were evaluated using an ethanol-induced ulcer model [51]. The methanol extracts was given at dosages of 250 and 500 milligram/kg. Ranitidine concentration was used as the standard. Pre-treating the extract led to a statistically significant decrease in both the free and total acidity of the stomach fluid. Only at a higher dosage of 500 mg/kg was there a significantly increased likelihood of the gastric juice pH altering. The methanolic extract of A. indicum exhibited potent anti-ulcer effects, which were influenced by the dose administered.

Analgesic and sedative effects

An experiment was conducted to study the pain-relieving and calming effects of a hydroalcoholic extract from the upper parts of A. indicum on albino mice [52]. Various pain tests were used, such as the eddy's hotplate test, the acetic acid induced writhing test, the hot immersion test, the tail clip test, and the actophotometer test to assess sedative effects. The extract was administered at a dosage of 400 mg/kg. The extract's significant analgesic and sedative effects are believed to be caused by opening the chloride channel or closing the sodium and/or calcium channels.

Diuretic function

The study examined the diuretic effects of ethanol, methanol, and ethyl acetate extracts from the leaves of A. indicum and Amaranthus spinosus on albino mice. The extract dose used varied from 100 to 400 mg/kg. Furosemide was regarded as the benchmark in the pharmaceutical sector. The ethanolic and methanolic extracts exhibited potent diuretic effects, but the ethylacetate fraction showed little diuretic activity [53].

Antifungal properties

The antifungal properties of the methanolic extract from different portions of A. indicum were evaluated against 11 fungal strains. The fungal strains tested were A. flavus, A. niger, A. fumigatus, C. albicans, C. utilis, Fusarium oxysporum, F. solani, Microsporum gypsum, Trichophyton metagraphytes, and Epidermophyton flocosum. Ketoconazole was used as the reference. The antifungal properties of the methanolic leaf extract of A. indicum were much greater than those of Trichophyton rubrum [54].

Larvicidal action

The study examined the larvicidal effects of extracts from 5 medicinal herbs, namely A. indicum, Aegle marmelos, Euphorbia thymifolia, Jatropha gossypifolia, and Solanum torvum, on the larvae of C. quinquefasicatus. The plants were analyzed in terms of total larvicidal efficacy. The pet-ether extract of A. indicum exhibited the highest larval death rate compared to the other extracts [55]. The bioassay of A. indicum identified and separated beta sitosterol, a potential mosquito larvicide having LC50 values of 11.49, 3.58, and 26.67 ppm against Aedes aegypti, Anopheles stephensi, and C.quiquefasciatus, respectively.





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Anti-diabetic effect

The study examined the antidiabetic activity of the chloroform fraction of the A. indicum plant extract on streptozocin-induced diabetic male Sprague Dawley rats. Analyses were conducted on glycosylated hemoglobin, total hemoglobin, plasma insulin, and glucose levels. Diabetic rats treated with chloroform fraction showed a substantial decrease in blood sugar level [56].

Anticonvulsant efficacy

The anticonvulsant properties of A. indicum leaves were examined in male Wistar rats that were subjected to either maximum electrical shock (MES) or pentylene tetrazole (PTZ)-induced convulsions [57]. Observations were conducted on the time of death, seizure latency, and fatality rate. Diazepam was the standard medication that was given. The plant extract doses provided were 100 and 400 mg/kg. The extracts showed anticonvulsant effects in the PTZ paradigm by prolonging latency and reducing the onset of clonic convulsions and clonic seizure activity. When MES was employed to induce convulsions, the extracts exhibited an anticonvulsant action by prolonging the latency to clonic extension and reducing the duration of tonic extension. An extract from A. indicum showed substantial anti-epileptic effects against several epileptic medications.

Anti-diarrheal effect

The anti-diarrheal effect of pet-ether and normal water extract of A. indicum leaf was treated in Wistar albino rats using methods such as GI motility, castor oil induced diarrhea, and prostaglandin E2-induced enter pooling [58].. The given dosage of the extracts was 500 mg/kg. The medicine of choice for therapy was loperamide. The A. indicum extract decreased intestinal movement in animal models fed charcoal meal and significantly blocked PGE2-induced enteropooling. The nonspecific spasmolytic activity of A. indicum may be responsible for suppressing motility, which is its mechanism of action. The leaf extracts significantly decreased the amount and frequency of feces excretion, similar to the medication loperamide. Both the methanol and aqueous extracts demonstrated anti-diarrheal effectiveness in PGE2- and castor oil-induced models of diarrhea, respectively. A. indicum exhibits considerably greater anti-diarrheal activity compared to loperamide.

Aphrodisiac effect

Rats were used to examine the aphrodisiac effects of water and ethanol extracts from the roots of A. indicum. An analysis of mating behavior, assessment of mating performance, study of hormones, and inspection of reproductive organs and sperm were conducted. The animals were administered at doses of 200 and 400 milligram per kilograms. Sildenafil citrate was used as a reference standard. The ethanolic extract exhibited potent aphrodisiac effects in male rats when administered at higher dosages (400 mg/kg). This was evidenced by a rise in the quantity of mounts, enhanced mating proficiency, and hormonal assessments. Both the ethanolic extract at a lower dose and the aqueous extract exhibited aphrodisiac effects. The research indicates that the root of A. indicum possesses aphrodisiac qualities [59].

Antioxidant function

The antioxidant capacity of organic solvent extracts from A. indicum and A. muticum was evaluated [60]. This experiment examined total phenolic content, total flavonoid content, DPPH free radical scavenging effects using ABTS+ decolorization test and FRAP assay to determine Trolox equivalent antioxidant capacity, and lipid peroxidation value. Total phenolic compounds were determined using the technique established by Singleton and Rossi (1965). Colorimetry was employed to determine the overall flavonoid content. The antioxidant levels in the above-ground parts were higher than those in the roots. The ethyl acetate fraction has the highest amount of phenolics. The ethyl acetate extract of A. muticum aerial parts and the butanol extract of A. muticum roots had the highest TEAC values among all samples, indicating substantial ABTS radical scavenging action. Both plant species shown strong free radical scavenging abilities when their DPPH radical scavenging effects were assessed using the method suggested by Sanchez-Moreno et al. (1998). The reduction capacity was assessed using an adapted version of the FRAP test established by Benzie and Strain (1960). The experiment's results suggest that antioxidant activity rises in correlation with the divergence of the solvent utilized for extraction. Ferric thiocyanate is utilized in an emulsion





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system including linoleic acid to determine the lipid peroxidation value. The value generated by the root extracts was comparable to that of the gold standard antioxidant, Trolox. This thorough investigation indicates that both species possess a heightened antioxidant capacity, which might potentially be used for addressing a range of concerns associated with degenerative diseases [61].

Immunomodulatory effects

In this investigation, albino mice were used to examine the immunomodulatory effects of an ethanolic and an aqueous extract of A. indicum leaves [62]. The doses for both extracts were 200 and 400 mg/kg/day, given orally, respectively. Hemagglutination antibody titers, delayed type of hypersensitivity, neutrophil adhesion, and carbon clearance tests all verified the immunomodulatory function. Oral administration of A. indicum extracts significantly increased antibody production in response to sheep red blood cells, according to the study's results. This study found that heamagglutination antibody titres increased considerably in both the primary and secondary groups, but in the cyclophosphamide-treated group, A. indicum exhibited a much higher titre. The delayed kind of hypersensitive reaction was significantly amplified in mice by A. indicum, which facilitated the footpad thickness response to sheep red blood cells. The proportion of neutrophils adhering to nylon fibers and their phagocytic activity were both significantly increased by A. indicum.

CONCLUSION

A. indicum has a wide variety of pharmacological effects, including hepatoprotective, wound-healing, immunomodulatory, analgesic, antibacterial, antimalarial, and hypoglycemic effects. Phytochemicals, glycosides, flavonoids, carbohydrates, tannins, and steroids constitute the majority of the plant's chemical composition. In an effort to help the public better understand A. indicum and its potential uses in alternative medicine, this review paper compiles and synthesizes relevant information.

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Table 1. Reported elements in the different parts of A. indicum [27]

Plant parts/									
Elements (p.p.m)	Na	K	Ca	Mg	Fe	Cu	Co	Mn	Ni
Leaves	486	11854	117241	55862	3190	109	38	326	52
Flowers	390	14088	103898	44746	1990	90	35	325	52
Seeds	567	6788	33000	70400	1696	234	30	336	66
Stems	631	3782	85227	74091	2400	133	47	159	61
Roots	2820	3936	41148	18899	2050	86	27	84	36

Table 2. Presenting various phytochemical constituents in the different parts of plant A. indicum

Phytochemical compounds	Parts	References
Vanillic		
p-coumaric		
p-hydroxybenzoic		
Caffeic and fumaric acids		
_Q -β-D-glucosyloxybenzoic	Aerial portion	[63]
Gluco-vanilloyl glucose	Actial portion	[65]





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4-(3-Hydroxybutyl) phenol		
4H-Pyran-4-one, 2,3-dihydro-3,5-di hydroxy-6-methyl-		
2-Methoxy-4-vinylphenol		
Phenol, 2,6-dimethoxy-		
Formic acid, 2,6-dimethoxyphenyl ester		
Ethanone, 1-(2-hydroxy-5-methylphenyl)-		
Pyrazole, 4-ethyl-3,5-dipropyl-		
2-(1-Methyl-2-propenyl)bicyclo[2.2.1]heptane		
1H-Pyrrole, 2-(2,4,6-cycloheptatrienyl)		
4-((1E)-3-Hydroxy-1-propenyl)-2-methoxyphenol		
2-Propenoic acid, 3-(4-hydroxy-3-methoxyphenyl)-, methyl ester		
Fatty acid (palmitic)		
Oxacyclododecane-2,8-dione		
Ethyl 9,12,15-octadecatrienoate		
Fatty acid (linoleic)	Aerial portion	[64]
Phenol, 4,4'-(1-methylethylidene)bis-	7 ichai portion	[O±]
N-(2-(4-hydroxyphenyl)ethyl] acetamide		
2,6,10,14,18,22-Tetracosahexaen (squalene-triterpene)		
4α-Methyl-1-methylidene-1,2,3,4,4a,9,10, 10a-Octahydrophenanthrene	1	
Piperine		
aurantiamide acetate		
Methyl indole-3-carboxylate		
3,7-dihydroxychromen-2-one		
scoparone		
scopoletin	=	
syringaldehyde	1	
1-methoxycarbonyl-b-carboline	1	
1-lycoperodine		
3-hydroxy-beta-damascone	=	
adenosine		
p-hydroxybenzoic acid		
3-hydroxy-b-ionol		
N-feruloyl tyrosine	Whole plants	
Abutilin A (1)	1	
(R)-N-(10-Methoxycarbonyl-20-phenylethyl)-4-hydroxybenzamide	1	[65]
Beta-sitosterol	1	[00]
stigmasterol	1	
vanillin		
methylcoumarate	-	
4-hydroxyacetophenone	-	
p-hydroxybenzaldehyde	-	
vanillic acid	-	
Benzoic acid	1	
methylparaben	1	
71	1	
4-hydroxy-3-methoxy-		
trans-cinnamic acid methyl ester	1	
trans-p-coumaric acid	-	
thymine		





adenine		
methyl 4-hydroxyphenylacetate		
riboflavin		
p-b-D-Glucosyloxybenzoic acid		
p-Hydroxybenzoic		
Caffeic acid	Whole plants	[66]

Table 3. Presenting various phytochemical constituents in the root of plant A. indicum

Phytochemical compounds	References
Proteins	
Free Acid	
Resin	
Linoleic	
Stearic	
Palmitic	
Lauric	
Myristic	
Caprylic,	
Capric	
Sitosterol,	
Abutilin A	[67]
(R)-N-(1'-methoxycarbonyl-2'-	
phenylethyl)-4-hydroxybenzamide	
Carbohydrates	
Free amino acids	[68]
Nonanoic acid	
Undecanoic acid	
Hexadecanoic acid, methyl ester	
N-Hexadecanoic acid	
Methyl 8,9-methylene-heptadec-8-enoate	
Trichloroacetic acid, dodec-9-ynyl ester	
9,12-octadecadienoic acid (Z, Z)-	
Squalene	
Campesterol	[69]
Stigmasterol	[02]
Beta-sitosterol	

Table 4. Presenting various phytochemical constituents in the leaves of plant A. indicum

Phytochemical compounds	References
Alkaloids	[2]
Amino acid	[62]
Mucilage	
Organic acid	
Triterpenoids	
glycosides	
Phenols	[70]
Steroids	





N-Acetyl-N-desmethylmethoxyphenamine	
Diphenylmethane	
Cathinone	
Methyl-alpha-d-ribofuranoside	
3,7,11,15-tetramethyl-2-hexadecen-1-OL	
7,9-di-tert-butyl-1-oxaspiro (4,5) deca-6,9-diene-2,8-dione	
1,2-benzenedicarboxylic acid, butyl octyl ester	
1,3,6-heptatriene, 2,5,5-trimethyl	
Benzene, 1,4-bis (phenylmethyl)-	[71]
Octadecane, 9-ethyl-9-heptyl	
Arabinitolpentaacetate	
Hydroxybenzoic acid ester	
3 hydroxy beta ionol	
Abutilin A	
Z-11-Hexadecenoicacid	
4-Methylcholestan-3-ol-,(3 β ,4 α ,5 α)-;4- α -Methyl-5- α -cholestan-3- β -ol	
10-Hydroxy-2 decenoic acid methyl ester	
[1,1'-bicyclopropyl]-2-octanoic acid, 2'-hexyl-, methyl ester	
4 hydroxyphenylacetic acid methyl ester	
5-Thio-D-glucose	[72]
(E)-10-Heptadecen-8-ynoicacid methyl ester	[72]
5-Allylsulfanyl-1-(4-methoxy-phenyl)-1H-tetrazole	
9,10-Anthracenedione,1,4diamino-2-methoxy	
Triamcinolone Acetonid	
10-Methoxydihydrocorynantheol;10-methoxycorynan-17-ol	
(R)-N-(1'-methoxycarbonyl-2'-phenylethyl)-4- hydroxybenzamide	[72, 73]
phytol	[71]
Ergostanol	[73]
Carbohydrates	[2]
Proteins	
Phenols	
Saponins	5643
Steroids	[61]
Reducing sugars	
Tannins	103
Flavonoids	[2]
L	

Table 5. Presenting various phytochemical constituents in the flowers of plant A. indicum

Phytochemical compounds	References
Methylstigmasterol	
Triacontanoic acid	
Apigenin 7-0-beta-rhamnopyranosyl	
Uresenol	
Glycopyronoside	
Luteolin	
Chrysoeriol -7-0-beta -glucopyranoside	
Quercetin 3-0-alpha – rhamnopyranosyl(1-6)-beta-glucopyranoside	





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Quercetin	
Apigenin	
Chrysoeriol	
Glucopyranoside	[33-36, 70]
7-0-beta glucopyranoside	
Quercetin 7-0-beta-glucopyranoside	
Abutlin	
(R)-N-(1-methoxycarbonyl-2-phenylethyl)-4-hydroxy benzamide	

Table 6. Presenting various phytochemical constituents in the fruits of plant A. indicum

Phytochemical compounds	Reference
2-Pentanone, 4-hydroxy-4methyl-,	
2-Hexanol, 2-methyl	
2-Pentanol, 2,3dimethyl	
m-xylene,	
p-xylene	
o-xylene,	
c-Sitosterol,	
a-Sitosterol	
Cholest-5-en-3-ol	
4,4-dimethyl-,(3a)-	[37, 38]
Lupeol, Lup-20(29)-en-3ol,acetate,(3a)	[57,50]
9,19-Cyclo-9a-lanostane3a,25-diol	
Taraxasterol	

Table 7. Presenting various phytochemical constituents in the seeds of plant A. indicum

Phytochemical compounds	References
The seed contains protein. The amino acid composition of protein is: proline, glycine, alanine, cysteine,	
aspargine, threonine, serine, glutamine, methionine, isoleucine, valine, leucine, tyrosine,	
phenylalanine, histidine, lysine and arginine.	[39]
Galactomannose	
D-galactose	[70]
D-mannose	[70]

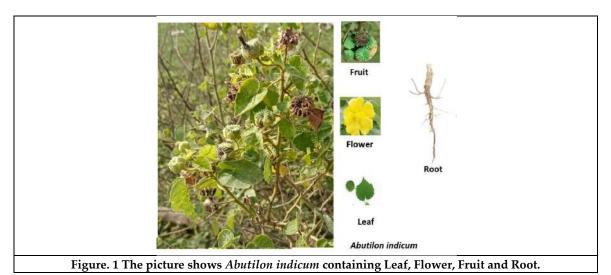
Table 8. Presenting various pharmacological activities of A. indicum

Sl. No.	Activity	References
1.	Antibacterial	[41]
2.	Anti-inflammatory	[42]
3.	Hepatoprotective	[43]
4.	Nephroprotective	[44]
5.	Wound healing	[45]
6.	Anti-venom	[46]
7.	Anticancer	[47]
8.	Anti-arthritic	[48]
9.	Anti-Alzheimer's	[49]
10.	Hypoglycemic	[50]
11.	Anti-Ulcer	[51]





12.	Analgesic & Sedative	[52]
13.	Diuretic	[53]
14.	Anti fungal	[54]
15.	Larvicidal	[55]
16.	Anti-diabetic	[56]
17.	Anti-convulsant	[57]
18.	Anti -diarrheal	[58]
19.	Aphrodisiac	[59]
20.	Antioxidant	[60, 61]
21.	Immunomodulatory	[62]







RESEARCH ARTICLE

Tourism Sustainability at Mahabalipuram Heritage Site - Impact on **Farmer Community**

P. Veeralakshmi¹, Prabayathi Venugopal², Jeevanantham. N³ and M. Sampath Nagi^{4*}

- ¹Assistant Professor, Department of Commerce, Government Arts and Science College, Kallakurichi, Tamil Nadu, India
- ²Associate Professor, School of Management Studies, Jai Shriram Engineering College, Tirupur, (Affiliated to Anna University, Chennai), Tamil Nadu, India
- ³Assistant Professor, Department of Business Administration, Government Arts and Science College, Valparai, (Affiliated to Bharathiyar University), Coimbatore, Tamil Nadu, India
- ⁴Assistant Professor, Department of Management Studies, Pondicherry University, Port Blair Campus, Pondicherry, India

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*Address for Correspondence

M. Sampath Nagi

Assistant Professor, Department of Management Studies, Pondicherry University, Port Blair Campus, Pondicherry, India Email: sampathnagi@gmail.com



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ABSTRACT

Tourism has today achieved the status of a complete industry today. "Travel and tourism enhance the quality of human experience in spiritual as well as material way". The tourism provides employment to a wide spectrum of job seekers from the unskilled to the specialized. Heritage is regarded as one of the most significant and fastest growing components of tourism in many developed economies. Considering the importance of residents who have a key role in tourism, many studies have been carried out by researchers in developed countries about local farmer community perceptions toward tourism. But unfortunately, hardly any work has been devoted to examine the residents' perception on the impact on tourism sustainability development in developing countries. The research for the current study includes descriptive research. The data collected for the research consist of primary and secondary data. This research has an applied questionnaire as the research instrument for collecting the data. The questionnaire has four factors like social, cultural, economic & environmental aspects and major dimension as tourism sustainability factors enhancing local farmer community The researcher has considered a sample of 384, with a confidence level of 95% and a margin of errors of 5% from the Demorgan's sample size estimator table. The statistical tools used for analysis are measures of central tendency and dispersion, bivariate correlation and structural equation modeling. The results propose that





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while tourism practices have been successful in positively affecting the economy, there is room for improvement in addressing the negative impacts on social, cultural, and environmental factors. Efforts should be directed toward creating a more balanced and sustainable approach to tourism that takes into account the well-being of the local farmer community, the preservation of cultural heritage and the protection of the environment.

Keywords: Tourism, Heritage, Sustainability, Local farmer Community, Social, Cultural, Economic & Environmental.

INTRODUCTION

Tourism has today achieved the status of a complete industry today. "Travel and tourism enhance the quality of human experience in spiritual as well as material way" as quoted by Krippendorf, 1987 [1], Lane (2018) [2]. For many developing countries, it is one of the main sources of foreign exchange income and the number one export category, creating the much-needed employment and opportunities for development. As a highly labour-intensive activity, tourism and tourism support activities create a high proportion of employment and career opportunities for low skilled and semi-skilled workers, particularly for the poor, female and young workers. In India, the travel and tourism sector are estimated to create 78 jobs per million rupees of investment compared to 45 jobs in the manufacturing sector for similar investment. Apart from providing employment to a wide spectrum of job seekers from the unskilled to the specialized, a higher proportion of tourism benefits (jobs, petty trade opportunities) accrue to women.

Heritage Tourism

Heritage is regarded as one of the most significant and fastest growing components of tourism in many developed economies as per the authors (Chauhan. (2022) [3], Alzue & Morrison (1998) [4]; Herbert (2001) [5]. It is becoming increasingly popular in the world, and deemed to be important for tourism development. Defined as a form of special interest tourism, it caters to the desire of tourists interested in learning about the history and lifestyle of a destination (Li & Lo (2005) [6], Craik (1997) [7], Williams & Lawson (2001) [8]. It has long been recognized that the ideological and institutional context of heritage tourism is fundamentally different from that of general tourism, recited by Garrod & Fyall (2000) [9]. Tourism is widely acknowledged for its role in creating national and / or regional wealth Phrased in UNWTO (2009) [10]. The authors Stronza (2007) [11]; Weaver & Lawton (2007) [12] said, if sustainably managed (Sampath et.al, 2022) [13], tourism can provide numerous other benefits. Tourism can be a catalyst for the conservation and revival of natural and cultural heritage and for the promotion of the positive image of people and places. Tourism can enhance global awareness of the cultural and natural values of destinations, especially those in developing countries (Henderson, 2003 [14]; Robinson & Picard, 2006) [15].

World Heritage Sites

According to UNESCO, the World Heritage Site is a site such as a forest, mountain, lake, desert, monument, building, complex, or city that is on the list and is maintained by the international World Heritage Programme administered by the UNESCO World Heritage Committee, composed of 21 state parties which is elected by the General Assembly for a four-year term. As of 2009, 890 sites in 148 countries are listed: 689 cultural, 176 natural, and 25 mixed properties in 148 states. Italy is home to the greatest number of World Heritage Sites to date with 44 sites on the list. The World Heritage Committee has divided the countries into five geographic zones: Africa, Arab States [composed of North Africa and the Middle East), Asia-Pacific (includes Australia and Oceania), Europe and the Americas (includes North America and South America). Russia and the Caucasus states are classified as European, while Mexico is classified as belonging to the Latin America and Caribbean zone. To foster heritage conservation, the World Heritage Convention (UNESCO, 2010) [16] has recognized, as of June 2010, 911 outstanding cultural and natural properties. Of these, 704 are cultural, 180 are natural, and 27 are mixed. Because of





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the outstanding cultural and conservation value of these properties, national and global efforts are being exerted to conserve and protect them. The World Tourism Organization (UNWTO, 2004) [17] has set the guidelines for sustainable tourism development and sustainable management practices based on social, economic and ecological aspects. These guidelines focus on making an optimal use of environmental resources, respecting the socio-cultural authenticity of host communities and ensuring viable, long-term economic operations (Nagi & Kumar, 2021) [18], (UNWTO, 2004) [15] a way of development intended to improve the quality of life of host communities, provide a top-quality experience for the visitors and preserve the quality of the environment.

Heritage Tourism in India

India's historical heritage is also very significant as it consists of secular and religious monuments dedicated to various faiths. The artistic and cultural heritage of the country is viewed as a USP by promoters and destination designers. According to various estimates, heritage tourism accounts for more than 60% of the overall share of tourists coming to India. It is overwhelming to note that 80% of the foreign tourists are interested in visiting rich heritage sites like Delhi, Jaipur, Agra and Mamallapuram and so on. India is ranked 11th in the Asia Pacific region and 40th overall, moving up three places on the list of the world's attractive destinations. It is ranked the 14th best tourist destination for its natural resources and 24th for its cultural resources, with many *World Heritage Sites*, both natural and cultural, rich fauna, and strong creative industries (Nagi, Vijayakanthan & Arasuraja, 2022) [19] in the country.

Significance of the Study

Today multiplier effects of tourism are considerably common for the national economies. Tourism is a bilateral event which covers not only locals but also the visitors. None of the tourist would like to encounter the inappropriate behaviors of the locals even if the destination shines with the tremendous natural environmental and historical beauties. To better understand the aspects of heritage and history that are utilized and the application of sustainable concepts in Mamallapuram, this investigation focuses strongly on the perceptions of the stakeholders towards sustainable tourism. The community, as hosts to tourists, is vital in the visitor experience and research suggests that it is impossible to sustain tourism to a destination that is not supported by the local people (Ahn, Lee & Shafer 2002; [20] Twinning-Ward and Butler 2002 [21]; McCool, Moisey and Nickerson 2001) [22]. The research focuses specifically on the impact of socioeconomic, cultural and environmental of heritage tourism towards the local community at Mamallapuram. In order to ascertain how sustainability is perceived by the host views and whether those perceptions are conducive to increasing the sustainability of heritage tourism.

Problem of the Study

Early work on perceived impacts of tourism, which dates back to the 1960s, tended to focus on the economic and positive effects of tourism (Pizam, 1978) [22]. However, in the 1970s, the consequences of tourism were examined more critically by anthropologists and sociologists who emphasised negative socio-cultural impacts (De Kadt, 1979) [23]. The 1980s and 1990s have been characterised by a more balanced perspective, recently called sustainable tourism, where positive and negative effects. (Ap & Crompton, 1998) [23]. The economic impacts of tourism are usually perceived positively by the residents. Tourism acts as an export industry by generating new revenues from external sources. A host nation will gain foreign exchange, which will contribute to improve the nation's balance of payments (Gee et al, 1997) [24]. It decreases unemployment by creating new job opportunities (Sheldon and Var, 1984) [25]. Increasing demand for tourism encourages new infrastructure investment (Inskeep, 1991) [26] and communication and transportation possibilities (Milman and Pizam, 1988) [27]. Residents of tourist destination might have a better standard of living and higher income by tourism activities. However, if not well planned and controlled, tourism may lead to negative impacts or reduce the effectiveness of positive ones. The prices of goods and services might go up with the increased demand from foreign customers (Liu and Var, 1986) [28]. Tourism might cause a gradual change in a society's values, beliefs and cultural practices. Local residents feel this impact more heavily. By observing the tourists, local people might change their life style (dressing, eating, entertainment and recreational activities, and so forth). While this influence may be interpreted positively as an increase in the standard of living, it may also be considered negatively as an indication of acculturation (Brunt and Courtney, 1999 [29],





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Dogan 1989) [30]. Tourism can contribute to the revitalisation of arts, crafts and local culture and to the realisation of cultural identity and heritage. In order to attract more tourists, architectural and heritage sites are restored and protected (Inskeep, 1991 [25], Liu and Var, 1986 [31]). Moreover, many people of different cultures come together by means of tourism, facilitating the exchange of cultures (Brayley et al, 1990) [32]. In addition to its cultural impacts, tourism is perceived to contribute to changes in value systems, individual behaviour, family relations, collective lifestyle, moral conduct and community organisations (Ap and Crompton, 1998) [22]. These kinds of social impacts may be positive or negative. With the development of tourism in an area, there might be changes in social structure of the community. Basically, two different classes; a rich class which consists of businessmen and landowners, and a lower class which contains mostly immigrants might emerge in the community (De Kadt, 1979 [21], Dogan, 1987 [29]). It also modifies internal structure of the community by dividing it into those who have and have not a relationship with tourism or tourists (Sampath et.al, 2023)[33]. (Brunt and Courtney, 1999 [28]). Intense immigration from different cultures of people brings about social conflict in the area. Generally, impacts of tourism on women are perceived positively such as more freedom, more opportunities to work, increase self-worked and respect, better education, higher standards of living with higher family income. However, some argue that tourism distracts family structure and values, and also leads to increase in divorce rates and prostitution (Gee et al, 1997 [23]).

Tourism may lead to a decline in moral values; invokes use of alcohol and drugs; increases crime rates and tension in the community (Liu and Var, 1986 [30] Milman and Pizam, 1988 [26 [26]). Moreover, with the development of tourism, human relations are commercialised while the non-economic relations begin to lose their importance in the community (Dogan, 1989 [29]). In relatively small tourism resort towns, increased population and crowd especially in summer seasons cause noise, pollution and congestion. This limits the use of public areas such as parks, gardens and beaches as well as of local services by the residents, which sometimes result in negative attitudes towards tourists (Ross, 1992 [34]). Local communities are known as the key stakeholders in leisure and tourism management. Tourism has to be managed with the help and interest of all stakeholders in a given territory with a focus on local inhabitants (Guyer and Pollard, 1997 [35]). Considering the importance of residents who have a key role in tourism, many studies have been carried out by researchers in developed countries about local community perceptions toward tourism (Lankford and Howard 1994 [36], Williams and Lawson, 2001 [37], Nicholas, 2007 [38]). But unfortunately, hardly any work has been devoted to examine the residents' perception on the impact on tourism sustainability development in developing countries (Lepp, 2007 [39]).

REVIEW OF LITERATURE

Ahmed, Nassar (2023) [40] in the study "Influence of Cultural heritage tourism image on resident perceived impacts and their support: Evidence from Jammu and Kashmir, India" states that every destination's tourism growth is intrinsically related to local support, particularly during periods of tourism planning, crises, and hostile movements to industry. The territory of Jammu and Kashmir within India serves as the venue for the investigation, wherein tourism based on cultural heritage is still in its infancy. The results demonstrate that inhabitants' assessments of theeconomic, socio cultural and environmental | consequences are substantially and favourably related to CHT image, which ultimately influences community support towards CHT development. Furthermore, the conclusions of this investigation illustrate the relevance of CHT image in determining residents' opinion of tourism effects and support, which has received very limited consideration in the realm of cultural heritage. The research advances overall knowledge of how inhabitants' opinions are evolved in underdeveloped cultural heritage areas, while stressing the need for measures that are more location-based, adaptable, and resident-centred. Lekaota (2018) [41] has researched on "Impacts of World heritage sites on local communities in the Indian Ocean Region". The IORA (Indian Ocean Rim Association) is committed to various objectives which include fostering tourism and cultural exchanges in the region. IROA is also committed in promoting cultural heritage and involving the economic potential of heritage including World Heritage properties and sites. The United Nations Education, Science and Cultural Organization (UNESCO), has declared many areas as World Heritage sites along the Indian Ocean. The study used a qualitative literature review method to unpack IORA countries and their world heritage sites. The sampling comprised the IORA





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countries. Purposive sampling was used for the selection of IORA countries. The results show lack of local community involvement in the management of world heritage sites, and thus no benefits accrue to them. The countries also showed that local community's participation could indeed contribute to sustainability of World Heritage Sites. The researcher recommends that IROA countries should form intense collaborations in order work together to have a common framework for managing World Heritage Sites which could benefit the communities. The researcher also proposes a benefit sharing model which can benefit IORA countries in seeking to achieve the main aim of the effective conservation of heritage and culture.

Research Proposed Model

The theoretical framework is a model of logical relationship amongst the variable of sustainability of tourism Figure 2, the self-developed theoretical model consists of a set of tourism dimension's (*social*, *cultural*, economic & environmental aspects) impact on the local community Based on this model, the research objectives are framed and hypotheses are also formulated.

Research Objectives

The host plays an important role in determining the success of tourism development. Without local participation there is no sustainable tourism (Kariel, 1982 [42]). The objectives of this study are to understand the whether the impact of social, cultural, economic & environmental aspects on the local communities is positive or negative at Mamallapuram.

- 1. To study the different degrees of tourism dimensions on the local community;
- 2. To identify the of tourism dimension relationship with respect to the local community.
- 3. To investigate the of tourism dimension impact with respect to the local community.

RESEARCH METHODOLOGY

The research for the current study includes descriptive research. The data collected for the research consist of primary and secondary data. This research has an applied questionnaire as the research instrument for collecting the data. In the questionnaire social, cultural, economic aspects carries the positive termed questions & environmental aspects has negative termed questions. So the respondents received based on the five point Likert scale has been justified in the interpretation of measures of central tendency and dispersion. The pilot study was conducted among the sample of 50 respondents. The calculated overall reliability coefficient (Cronbach Alpha) has exceeded 0.8 and appears to be invariably high across the entire variables. The population of the study are the local community like unorganized sector workers, farmers, industrial workers, merchants and etc. since the definite population is not available as per the government record the population is considered as infinite. The researcher has considered a sample of 384, with a confidence level of 95% and a margin of errors of 5% from the Demorgan's sample size estimator table. Around 400 questionnaires were distributed and 391 responses has been received back and remaining 9 questionnaires were found to be biased. The Statistical Package for Social Sciences (SPSS) 20 version and Analyzing Momentum of Structures (AMOS) 18 version were used for analysis. The statistical tools used for analysis are measures of central tendency (mean) and measures of dispersion (standard deviation), bivariate correlation and structural equation modelling.

Analysis

Measures of Central Tendency and Dispersion

The dimensionality tourism sustainability factors enhancing local community has four major dimensions measuring the level of social, cultural, economic & environmental aspects. The Measures of Central Tendency (mean) and Measures of Dispersion (Standard Deviation) has been used to measure. The results are displayed below; The mean value ranges from 4.63 to 1.64. The sustainability tourism practices have only one positive factor economical aspect (μ = 3.35), which is also supported in the previous study done by Chauhan. (2022) **Error! Bookmark not defined.** This clearly signifies that the respondents are able to reach their economical standards like increase in life





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standard, consumption and expenditure, property value and able to relate new job opportunities and investments. Whereas the social (μ = 1.64), which is also supported in the previous study done by De Kadt (1979) . & Dogan (1987), cultural (μ = 1.69), which is also supported in the previous study done by Brunt and Courtney (1999) & Dogan (1989) & environmental (negative questions) (μ = 4.36), which is also supported in the previous study done by Ahn, Lee & Shafer 2002; Twinning-Ward and Butler 2002 . McCool, Moisey and Nickerson 2001. factor show the negativity part of the tourism. This shows that in these factors respondents are feeling that they are not able to reach their standards.

Bivariate Correlation

The relationship among the dimensionality tourism sustainability factors enhancing local community (social, cultural, economic & environmental aspects) has been measured using the Bivariate Correlation. The results are displayed below; The results of the bivariate table shows that social aspect has positive relationship with the cultural aspect (r = 0.943) & tourism sustainability factors (r = 0.655). Likewise, the cultural aspect has positive relationship with the tourism sustainability factors (r = 0.634). In the same way, the economical aspect has positive relationship with the environmental aspect (r = 0.561) & tourism sustainability factors (r = 0.253). And correspondingly environmental aspect has positive relationship with the tourism sustainability factors (r = 0.327). Whereas the cultural aspect has negative relationship with the environmental aspect (r = 0.140). In the same way, the social aspect has no relationship with economical aspect & environmental aspect and the cultural aspect has no relationship with the economical aspect. These results are supported by the previous authors research like Brunt and Courtney, 1999 . & Ross, 1992).

Structural Equation Modelling

The Impact of social, cultural, economic & environmental aspects over the dependent variable tourism sustainability factors enhancing local community has been measured using the Structural Equation Modelling. The results are displayed below; The GFI (Goodness of Fit) and AGFI (Adjusted Goodness of Fit Index) should be nearing to one or one indicates that the model is a good fit. In this model the GFI value is 0.956 and AGFI value is 0.971. This clearly implies that the model is a good fit. The regression weight table clearly shows that the social (estimate = 0.344), cultural (estimate = 0.262), economic (estimate = 0.257) & environmental (estimate = 0.071) aspects have positive impact on the tourism sustainability factors enhancing local community (De Kadt, 1979 [21]). The independent variable human resource interventions show 48.9 percentage of its variance when influenced by the dependent variables social, cultural, economic & environmental aspects (Ap & Crompton, 1998 [22]).

DISCUSSION AND CONCLUSION

The results propose that while tourism practices have been successful in positively affecting the economy, there is room for improvement in addressing the negative impacts on social, cultural, and environmental factors. Efforts should be directed toward creating a more balanced and sustainable approach to tourism that takes into account the well-being of the local community, the preservation of cultural heritage, and the protection of the environment, (Chen, 2023, provides the guidelines to the relevant authorities and regulators in developing and implementing the regulators regarding the sustainability of tourism growth by promoting economic and environmental conditions and effective tourism policies in the country [42]). The consistently lower mean values in the social, cultural, and environmental aspects suggest that respondents perceive these dimensions of tourism to fall short in meeting their standards. This indicates a negative perception of the impact of tourism on these aspects and calls for greater attention and improvement in sustainable tourism practices. In conclusion, the study reveals a mixed picture of sustainability in tourism, with positive economic outcomes but challenges (Nagi, Kannan & Ramasubramaniam, 2021) [43] in social, cultural, and environmental dimensions (Banga et. al, 2022) [44]. To achieve more comprehensive sustainability, it is crucial for stakeholders in the tourism industry to address the identified concerns and work towards a more balanced and responsible approach that takes into account the well-being of local





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communities, cultural heritage preservation, and environmental conservation. Further research and concerted efforts are needed to foster sustainable tourism practices that bring positive outcomes across all dimension.

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Table 1. Classification of the Heritage Sites

Zone	Natural Heritage	Cultural Heritage	Mixed Heritage	Total
Africa	33	42	3	78
Arab States	4	60	1	65
Asia-Pacific	48	129	9	186
Europe & North America	56	375	9	440
Latin America & Caribbean	35	83	3	121
Total	176	689	25	890

^{*} Secondary Data - Sites Configured in the Study: (Information is partially retrieved from the sites given in the bibliography)

Values are the number of Heritage sites present in Location

Table 2. Mean & Standard Deviation Tourism Sustainability Factors enhancing Local Community

Variables	Mean	Sd
Social Aspect	1.64	0.63
Cultural Aspect	1.69	0.62
Economical Aspect	3.35	0.48
Environmental Aspect	4.63	0.48
Tourism Sustainability Factors enhancing Local Community	2.83	0.55

Table 3. Ho: No significant relation between the dimensionality tourism sustainability factors enhancing local

community (social, cultural, economic & environmental aspects)

Variables	Social Aspect	Cultural Aspect	Economical Aspect	Environmental Aspect	Tourism Sustainability Factors		
Social Aspect	1						
Cultural Aspect	.943**	1					
Economical Aspect	031	043	1				
Environmental Aspect	084	140**	.561**	1			
Tourism Sustainability Factors	.655**	.634**	.253**	.327**	1		
**. Correlation is significant at the 0.01 level (2-tailed).							

Values in the table are Pearson Correlation values

Table 4. Impact of Social, Cultural, Economic & Environmental Aspects → Tourism Sustainability Factors Enhancing Local Community

Test for Model Fit	Values
GFI (Goodness of Fit)	0.956
AGFI (Adjusted Goodness of Fit)	0.971

Table 5. Impact of Social, Cultural, Economic & Environmental Aspects → Tourism Sustainability Factors Enhancing Local Community

Variable	Variable	UE	SE	S.E	C.R	P	
Tourism Sustainability Factors Enhancing Local Community	←	Environmental Aspects	0.344	0.409	0.03	11.289	***



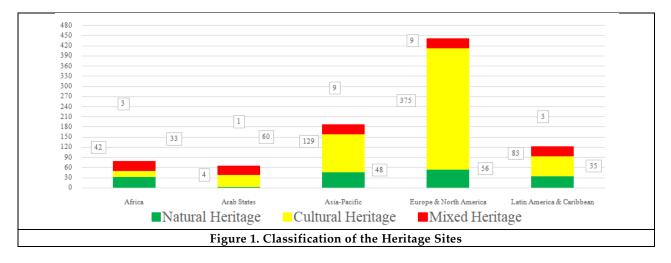


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Tourism Sustainability Factors Enhancing Local Community	←	Cultural Aspects	0.262		0.397	0.024	10.971	***
Tourism Sustainability Factors Enhancing Local Community	+	Social Aspects	0.257		0.397	0.023	10.968	***
Tourism Sustainability Factors Enhancing Local Community	+	Economic Aspects	0.071		0.083	0.031	2.301	0.021**
**Significant at 0.001 percent	age level			S.E - Standard Error				
*Significant at 0.05 percentage level				C.R - Critical Ration				
UE - Unstandardised Estimate				P - Probability Value				
SE- Standardised Estimate				Inf. – Influence				

Table 6. Impact of Social, Cultural, Economic & Environmental Aspects → Tourism Sustainability Factors Enhancing Local Community

Variable	Estimate
Tourism Sustainability Factors Enhancing Local Community	48.9







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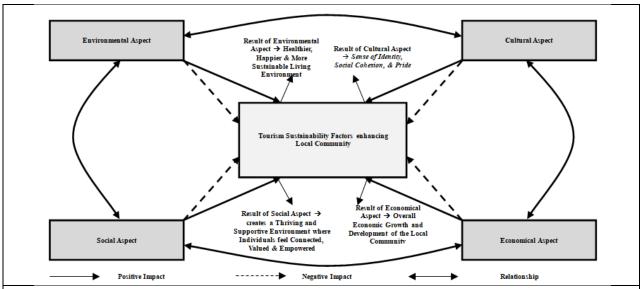


Figure 2. Tourism at Mahabalipuram Heritage Site - Impact on Local Community

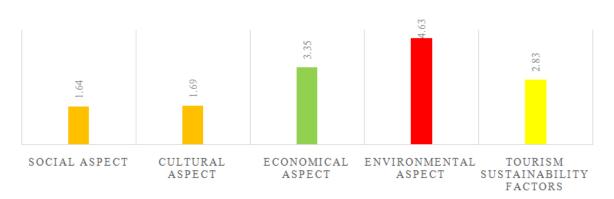


Figure 3. Tourism Sustainability Factors enhancing Local Community

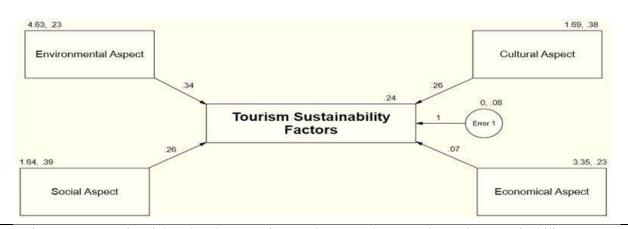


Figure 4. Impact of Social, Cultural, Economic & Environmental Aspects → Tourism Sustainability Factors
Enhancing Local Community





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RESEARCH ARTICLE

Analysis of Fuzzy Random Graphs for Decision Making

N. Sarala¹ and R. Abirami^{2*}

¹Associate Professor, Department of Mathematics, Department of Mathematics, A.D.M. College for Women (Autonomous), Nagappatinam (Affiliated to Bharathidasan University, Tiruchirappalli), Tamil Nadu, India.

²Research Scholar, Department of Mathematics, A.D.M. College for Women (Autonomous), Nagappatinam (Affiliated to Bharathidasan University, Tiruchirappalli), Tamil Nadu, India.

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*Address for Correspondence

R. Abirami

Research Scholar,

Department of Mathematics,

A.D.M. College for Women (Autonomous), Nagappatinam (Affiliated to Bharathidasan University, Tiruchirappalli),

Tamil Nadu, India.

Email: kannahiamf@gmail.com



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ABSTRACT

A key task for ensuring the effective running of an academic organization is the selection of competent and highly qualified academic staff. Due to the selection process' complexity, it is necessary to adopt the decision-making procedure, which allows for the most effective choice to be made based on the qualitative and quantitative information that is available. The primary focus of the current work is the selection of academic staff for an academic organization based on the vertex degree in the union and intersection of two fuzzy random graphs.

Keywords: Fuzzy Random Graph, Degree of vertex, characteristics of academic staff, Union and intersection of fuzzy random graphs.

INTRODUCTION

Employee dedication to the organization is what ensures its existence in nowadays aggressive competition. The success of an academic organization will indeed result from selecting potential hires who meet the requirements or policies to carry out a specific task. In literature, personnel selection receives a lot of attention. The Proceedings of the International Conference on Economics, Management, and Accounting (ICEMAC 2021) describe the proposed fuzzy logic for an intelligent decision support system to evaluate a student's eligibility for admission to the university. The applicability of the graph theoretic technique is first presented in 2016 to rank and choose academic staff out of ten





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candidates. A survey will be conducted in 2013 to examine the current fuzzy strategies for handling the staff selection process. It organizes various research techniques already in use and highlights gaps that require more investigation. Along with several fuzzy variations of fundamental concepts from graph theory, such as path, cycle, connectedness, etc., Rosenfeld developed the idea of the fuzzy graph in 1975. The idea of fuzzy relations was first suggested by Zadeh in 1987. Mordeson defined the operations such as union, Cartesian product, and composition of two fuzzy graphs. NN and Peng.C.S. The concept of a random graph was first proposed by Paul Erdos and Alfred Renyi in 1959, expanding on the probabilistic technique to determine the existence of particular graph features. The use of possibility theory in Professor Lotfi Zadeh's fuzzy sets and fuzzy logic theory dates back to 1978. Didier Dubois and Henri Prade also made contributions to its development. This research uses the concept of the introduction of fuzzy random graphs to establish the crucial factors for selecting the academic staff for arts and science colleges.

PRELIMINARIES

Definition:2.1

A Fuzzy Graph denoted by $\mathbb{G} = (V, E)$ on the graph $\mathbb{G}^* = (V^*, E^*)$ is a pair of functions (σ, μ) where $\sigma: V \to [0,1]$ is a Fuzzy subset of a non-empty set V and $\mu: V \times V \to [0,1]$ is a symmetric Fuzzy relation on σ such that for all u and v, the relation $\mu(u, v) = \mu(uv) \le \sigma(u) \land \sigma(v)$ is satisfied.

Definition:2.2

A Fuzzy Graph $\mathbb{H} = (V, \tau, \varphi)$ is called a Partial Fuzzy Subgraph of $\mathbb{G} = (V, \sigma, \mu)$ if $\tau(u) \leq \sigma(u)$ for every vertex $u \in \sigma$ and $\varphi(uv) \leq \mu(uv)$ for every $uv \in \mu$. In particular, $\mathbb{H} = (V, \tau, \varphi)$ is called a Fuzzy subgraph of $\mathbb{G} = (V, \sigma, \mu)$ if $\tau(u) = \sigma(u)$ for every $u \in \tau$ and $\varphi(uv) = \mu(uv)$ for every $uv \in \varphi$.

Definition:2.3

A probability space of all unlabled graphs with n vertices and m edges is called an Erdos-Renyi Random graph G(n,m). Each graph in this space has equally likely probability of being chosen, that is $P(\mathbb{G}_{n,m} = \mathbb{G}) = \binom{n}{2}^{-1}$. This formula follows as we are choosing m edges out $\binom{n}{2}$ possible edges, where each graph occurs with equal probability.

Definition:2.4

A Gilbert Random Graph $\mathbb{G}(n, p)$ is a probability space on the set of graphs with n vertices where each edge in the graph is added independently with probability P.

Definition:2.5

Let $\mathbb{V} = \{v_i, i = 1, 2, ..., n\}$ be a set of n vertices has n(n-1)/2 possible edges \mathbb{E} between them. Then there are two sets of edges, $\mathbb{E}_{\mathcal{F}} = \{(u, v)/1 \le u < v \le n \text{ and } (u, v) \text{ are Fuzzy edges}\}$

 $\mathbb{E}_{\mathcal{R}} = \{(u,v)/\ 1 \leq u < v \leq n \ \text{and} \ (u,v) \ \text{are Random edges} \} \ \text{that are disjoint. Consider the mapping}$

$$\begin{split} \phi_{\mathcal{G}} \colon \mathbb{V} &\to [0,1] \qquad u \to \phi_{\mathcal{G}}(u) \\ \psi_{\mathcal{G}} \colon \mathbb{V} \times \mathbb{V} &\to [0,1]_P \times [0,1]_{f_A} \end{split}$$

$$(\mathbf{u}, \mathbf{v}) \to \psi_G(\mathbf{u}, \mathbf{v}) = (P(\mathbf{u}, \mathbf{v}), f_A(\mathbf{u}, \mathbf{v}))$$

 $(u,v) \to \psi_{\mathcal{G}}(u,v) = (P(u,v),f_A(u,v))$ with P(u,v) = 0 if and only if $f_A(u,v) = 0$. Then the quartet $\mathbb{G} = (\mathbb{V},\mathbb{E},\phi_{\mathcal{G}},\psi_{\mathcal{G}})$ is called a Fuzzy Random graph if $\psi_{G}(\mathbf{u}, \mathbf{v}) \leq \min \mathbb{E} \varphi_{G}(\mathbf{u}), \varphi_{G}(\mathbf{v})$.

where (u, v) corresponds to the edge between u and v, P(u, v) and $f_A(u, v)$ represents the probability of the edge (u, v)and the membership function for the edge (u, v) within the fuzzy set A in Xrespectively.





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Definition:2.6

A Fuzzy Random Graph $\mathbb{H} = (\mathbb{V}, \mathbb{E}, \tau_{\mathcal{G}}, \sigma_{\mathcal{G}})$ is called a Partial Fuzzy Random Subgraph of $\mathbb{G} = (\mathbb{V}, \mathbb{E}, \varphi_{\mathcal{G}}, \psi_{\mathcal{G}})$ if $\tau_{\mathcal{G}}(u) \, \leq \, \phi_{\mathcal{G}}(u) \text{ for every } v \, \in \, \mathbb{V} \text{ and } \, \sigma_{\mathcal{G}} \, (uv) \, \leq \, \psi_{\mathcal{G}}(uv) \text{ for every } \, u \, \in \, \mathbb{V}. \text{ Also the Fuzzy Random Graph } \mathbb{H} = 0$ $(\mathbb{V}, \mathbb{E}, \tau_g, \sigma_g)$ is called a Fuzzy Random Subgraph of \mathbb{G} induced by α if $\alpha \subseteq \mathbb{V}$, $\tau_g(u) = \varphi_g(u)$ for all $u \in \alpha$ and $\sigma_{\mathcal{G}}$ (uv) = $\psi_{\mathcal{G}}$ (uv) for all u, v $\in \mathbb{E}$. We write $\langle \alpha \rangle$ to denote the Fuzzy Random subgraph induced by α .

Definition:2.7

In a Fuzzy Random Graph $\mathbb{G} = (\mathbb{V}, \mathbb{E}, \varphi_c, \psi_c)$ the degree of a Fuzzy Random vertex u is defined as

$$\begin{split} d_{\mathbb{G}}(u_i) &= \sum_{u_i \neq v_i \in \mathbb{E}_{\mathcal{F}}} \psi_{\mathcal{G}}(u,v) + \sum_{u_i \neq v_i \in \mathbb{E}_{\mathcal{R}}} nP \\ d_{\mathbb{G}}(u_i) &= \sum_{u_i, v_i \in \mathbb{E}_{\mathcal{F}}} \psi_{\mathcal{G}}(u,v) + \sum_{u_i, v_i \in \mathbb{E}_{\mathcal{R}}} n\psi_{\mathcal{G}}(u,v) \end{split}$$

for $(u, v) \in \mathbb{E}$ and $\psi_{G}(u, v) = 0$ for (u, v) not in \mathbb{E} . Where $\psi_{G}(u, v) = P$

Definition:2.8

The Union of two Fuzzy Random Graphs \mathbb{G}_1 and \mathbb{G}_2 is defined as a Fuzzy Random graph $\mathbb{G}_3=\mathbb{G}_1\cup\mathbb{G}_2:(\phi_{\mathcal{G}_1}\cup\phi_{\mathcal{G}_2},\psi_{\mathcal{G}_1}\cup\psi_{\mathcal{G}_2})\text{, where }\mathbb{V}_3=\mathbb{V}_1\cup\mathbb{V}_2\text{ and }$ $\mathbb{E}_3 = \mathbb{E}_1 \cup \mathbb{E}_2$ with

The Intersection of two Fuzzy Random graphs \mathbb{G}_1 and \mathbb{G}_2 is defined as a Fuzzy Random graph $\mathbb{G} = \mathbb{G}_1 \cap \mathbb{G}_2$: $(\varphi_{c_*} \cap \mathbb{G}_2)$
$$\begin{split} &\phi_{\mathcal{G}_2}, \psi_{\mathcal{G}_1} \cap \psi_{\mathcal{G}_2}) \text{, where } \mathbb{V} = \mathbb{V}_1 \cap \mathbb{V}_2 \text{ and } \mathbb{E} = \mathbb{E}_1 \cap \mathbb{E}_2 \text{ with} \\ &\left(\phi_{\mathcal{G}_1} \cap \phi_{\mathcal{G}_2}\right) (u) = \begin{cases} \phi_{\mathcal{G}_1}(u) \wedge \phi_{\mathcal{G}_2}(u); \text{ if } u \in \mathbb{V}_1 \cap \mathbb{V}_2 \\ 0; \text{ Otherwise} \end{cases} \text{ and } \end{split}$$

$$(\psi_{\mathcal{G}_1} \cap \psi_{\mathcal{G}_2})(uv) = \begin{cases} \psi_{\mathcal{G}_1}(v) \wedge \psi_{\mathcal{G}_2}(v) \text{ if } uv \in \mathbb{E}_2 \cap \mathbb{E}_1 \\ 0; \text{ Otherwise} \end{cases}$$

DECISION MAKING ON ACADEMIC STAFF SELECTION

Expected Characteristics of academic staff:

In the Arts and Science Colleges, there is a high expectation for quality academic staff. There are several reasons for this expectation.

- Experience
- **Educational Qualification**
- Salary expectations
- Ability to handle different subjects





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- · Research activities
- Technical skills
- Communication skills

Algorithm

Step 1: Input the Fuzzy Random vertex V_R

Step 2: Input the Fuzzy Random edge E_R

Step 3: Create the Fuzzy Random graphs

Step 4: Combine two Fuzzy random graphs by finding their union and intersection

Step 5: Determine the degree of vertices in Fuzzy Random graphs

Step 6: Identify the maximum degree of vertices and conclude that this characteristic is important.

Step 7: If the maximum value exceeds one value, repeat step: 1 and reassess the academic staff's characteristics.

Structure of the Problem

Step: 1

Suppose that there are two sets A and B which are two different background of candidates as far as the location of their institution they previously studied, namely urban and rural. Three candidates C_1 , C_2 , C_3 were taken as parameters. The following characteristics,

- u₁ stands for Experience
- u₂ stands for Educational Qualification
- u₃ stands for Ability to handle different subjects
- u₄ stands for Research Activities
- us stands for Technical skills

which are to be expected from the candidates were taken as the vertices of a Fuzzy random graph.

Step: 2

```
 \begin{split} \text{Consider} \ \mathbb{V} &= \{u_1, u_2, u_3, u_4, u_5\} \ \text{ and } \mathbb{E} = \{u_1 u_2, u_2 u_3, u_3 u_4, u_4 u_5, u_5 u_1, u_2 u_5, u_1 u_3, u_2 u_4\} \\ \text{Let} \ \mathbb{V}_1 &= \{u_1, u_2, u_3, u_4\} \quad \mathbb{E}_{\mathcal{F}_1} = \{u_1 u_2, u_2 u_3, u_4 u_5\} \quad \mathbb{E}_{\mathcal{R}_1} = \{u_3 u_4, u_5 u_1\} \\ \mathbb{V}_2 &= \{u_1, u_2, u_3, u_4\} \quad \mathbb{E}_{\mathcal{F}_2} = \{u_1 u_2, u_2 u_3, u_4 u_5\} \quad \mathbb{E}_{\mathcal{R}_2} = \{u_2 u_4, u_5 u_1\} \\ \mathbb{V}_3 &= \{u_1, u_2, u_3, u_4\} \quad \mathbb{E}_{\mathcal{F}_3} = \{u_2 u_3, u_1 u_3\} \quad \mathbb{E}_{\mathcal{R}_3} = \{u_2 u_5, u_3 u_4\} \end{split}
```

Then \mathbb{G}_1 and \mathbb{G}_2 are defined by Tables.

Step: 4

The Union of two fuzzy random graphs \mathbb{G}_1 and \mathbb{G}_2 are given by Tables.

The intersection of two fuzzy random graphs \mathbb{G}_1 and \mathbb{G}_2 are given by Tables.

Step: 5

Degree of vertices in the union of two Fuzzy random graphs

Degree of vertices in the union of two Fuzzy random graphs

Step: 6

According to the preceding procedures, vertex \mathbf{u}_2 is more significant than the other vertices. In this case, too, the maximum value does not exceed one. As a result, we come to an end here. Because of this, the selection of academic staff at colleges of arts and sciences heavily considers educational qualifications.





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CONCLUSION

A conclusion was drawn in light of the findings above. Educational qualifications and the characteristics of academic staff play a significant role in how firms choose their employees. The flexibility of vertex degree in fuzzy random graphs is offered as a selection criterion for academic employees among three applicants.

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Table: 3.1

\mathbb{V}_1	u_1	$\mathbf{u_2}$	u_3	u_4	u_5	
C_1	0.9	0.6	0.5	0.8	0.4	
C_2	0.8	0.3	0.7	0.6	0.5	
C_3	0.7	0.5	0.4	0.9	0.6	

Table 3.2

\mathbb{E}_{1}	u_1u_2	u ₂ u ₃	u ₃ u ₄	u ₄ u ₅	u ₅ u ₁	u_2u_5	u_1u_3	u_2u_4
C_1	0.6	0.3	0.4	0.4	0.2	0	0	0
C_2	0.3	0.25	0	0.4	0.5	0	0	0.3
C_3	0	0.2	0.3	0	0	0.5	0.4	0

Table 3.3

\mathbb{V}_2	$\mathbf{u_1}$	$\mathbf{u_2}$	$\mathbf{u_3}$	$\mathbf{u_4}$	$\mathbf{u_5}$
$\mathbf{C_1}$	0.7	0.8	0.6	0.9	0.8
C_2	0.6	0.4	0.4	0.5	0.7
C_3	0.4	0.6	0.5	0.2	0.6





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Table 3.4

\mathbb{E}_2	u_1u_2	u_2u_3	u_3u_4	u ₄ u ₅	u_5u_1	u ₂ u ₅	u_1u_3	u_2u_4
C_1	0.7	0	0.5	0	0	0.8	0.5	0
C_2	0.4	0.3	0.35	0	0.6	0	0	0.4
C3	0	0.5	0.1	0.2	0.4	0.4	0.4	0

Table 3.5

$\mathbb{V}_1 \cup \mathbb{V}_2$	$\mathbf{u_1}$	$\mathbf{u_2}$	$\mathbf{u_3}$	u_4	$\mathbf{u_5}$
C_1	0.9	0.8	0.6	0.9	0.8
C_2	0.8	0.4	0.7	0.6	0.7
C_3	0.7	0.6	0.5	0.9	0.6

Table 3.6

$\mathbb{E}_1 \cup \mathbb{E}_2$	u_1u_2	u_2u_3	u_3u_4	u ₄ u ₅	u_5u_1	u_2u_5	u_1u_3	u ₂ u ₄
C_1	0.7	0.3	0.5	0.4	0.2	0.8	0.5	0
C ₂	0.4	0.3	0.35	0.4	0.6	0	0	0.4
C_3	0	0.5	0.3	0.2	0.4	0.5	0.4	0

Table 3.7

$\mathbb{V}_1 \cap \mathbb{V}_2$	u_1	$\mathbf{u_2}$	u_3	u_4	$\mathbf{u_5}$
C_1	0.7	0.6	0.5	0.8	0.4
C ₂	0.6	0.3	0.4	0.5	0.5
C_3	0.4	0.5	0.4	0.2	0.6

Table 3.8

$\mathbb{E}_1 \cap \mathbb{E}_2$	u_1u_2	u_2u_3	u_3u_4	u ₄ u ₅	u_5u_1	u ₂ u ₅	u_1u_3	u ₂ u ₄
C_1	0.6	0	0.4	0	0	0	0	0
C_2	0.3	0.25	0	0	0.5	0	0	0.3
C_3	0	0.2	0.1	0	0	0.4	0.4	0

Table 3.9

Degree of vertex in $\mathbb{G}_1 \cup \mathbb{G}_2$	$\mathbf{u_1}$	\mathbf{u}_2	$\mathbf{u_3}$	u_4	$\mathbf{u_5}$
C_1	2.3	2.5	1.8	1.4	1.6
C_2	1.6	1.5	0.65	1.55	1.6
C_3	0.8	1.5	1.5	0.8	1.6
Total	4.7	5.5	3.95	3.75	4.8

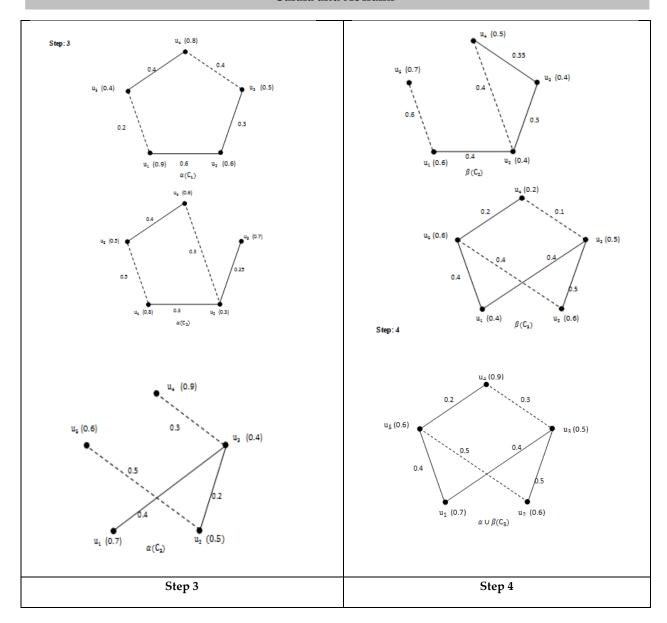
Table 3.10

Degree of vertex in $\mathbb{G}_1 \cap \mathbb{G}_2$	$\mathbf{u_1}$	$\mathbf{u_2}$	$\mathbf{u_3}$	$\mathbf{u_4}$	$\mathbf{u_5}$
C_1	0.6	0.6	0.8	0.8	0
C_2	1.3	1.15	0.25	0.6	1
C ₃	0.4	1	0.8	0.2	0.8
Total	2.3	2.75	1.85	1.6	1.8





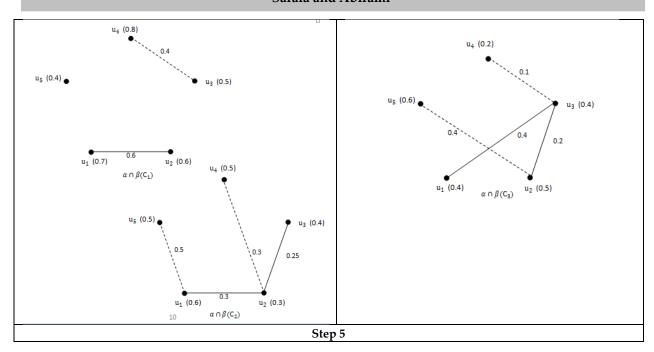
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RESEARCH ARTICLE

Assessing Periodontal Inflammation in Obese Individuals: Introducing Periodontal Inflammatory Surface Area (PISA) As A Comprehensive Measurement for Periodontitis -Pilot Study

V. Ramya¹ Julius Amaldas² J. Bhuvaneswarri³ and Aarthinisha⁴

- ¹Research Scholar; Professor Department of Periodontics, Sree Balaji Dental College, Biher University.
- ²Professor & HOD Department of Biochemistry, Sree Balaji Dental College.
- ⁴Senior Assistant Professor Department of Oral Medicine and Radiology, Taamilnadu Dental College and Hospital.
- ³Professor, Department of Periodontics, Sree Balaji Dental College.

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*Address for Correspondence

V. Ramya

Research Scholar & Professor, Department of Periodontics, Sree Balaji Dental College, Biher University.

Email: rammu82@gmail.com



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ABSTRACT

Recent research exploring the correlation between obesity and periodontitis has recognized obesity as a potential risk factor for the latter condition. Adipose tissue creates cytokines that promote inflammation, which might make periodontal tissues more inflamed. It's possible that conventional clinical measurements don't adequately reflect this influence. Digital pictures of periodontal pockets are used to assess inflammation accurately using the innovative Periodontal Inflammatory Surface Area approach, which was introduced in 2005. Due to the prevalent occurrence of obesity and periodontitis, scientists extensively examined the efficacy of PISA. This research underscores the necessity for improved diagnostic tools within this demographic and highlights the potential of PISA as a marker for inflammation. This study aims to assess PISA's usefulness in measuring inflammatory burden in periodontitis among obese individuals. A study was conducted at the Department of Periodontics, Sree Balaji Dental College & Hospital in Chennai, Tamil Nadu, involving twenty-one individuals categorized as obese or overweight. The recruitment followed the approval of the institutional human ethical committee. The study involved conducting a survey based on questionnaires to examine factors like diet, lifestyle, stress, body mass index, and waist-hip ratio. In evaluating periodontal health, measurements were taken for parameters such as bleeding on probing, recession, and clinical attachment. This data was





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then utilized to calculate Attachment Level Surface Area (ALSA) and Recession Surface Area (RSA). Subsequently, Periodontal Epithelial Surface Area (PESA) and Periodontal Inflammatory Surface Area (PISA) were derived through software analytic tools. The statistical analysis included the assessment of mean and standard deviation for continuous variables. A multivariate logistic regression analysis, conducted using STATA/SE13(Stata Corp., College Station, TX, USA) the software was designed to assess how risk factors influence the occurrence of periodontitis. Furthermore, Spearman's correlation analysis was conducted to investigate relationships between different variables. The study explored correlations between periodontal parameters, PESA/PISA, and obesity levels. Spearman's correlation analysis indicated a lack of significant correlation among BMI, BOP, age, waist measurement, and PISA score. In the investigated humanity, study showed no connection among PISA and obese status. However, higher BOP values correlated with elevated PISA scores.

Keywords: gingivitis, Infectobesity, PISA, and adiposity.

INTRODUCTION

Obesity is a prevalent concern on a global scale, especially in developed countries, as highlighted by the World Health Organization in 2006. [1]. This metabolic condition arises from an energy imbalance where calorie intake surpasses expenditure, resulting in a BMI exceeding 30.0 kg/m². Consequently, it leads to an accumulation of adipose tissue, as discussed by Bray in 2007. Among the prevalent chronic ailments, periodontal disease stands out. Periodontitis involves inflammation of the periodontal apparatus, leading to the deterioration of connective attachments and tooth loss. Certainly, obesity is a pervasive condition that impacts the onset and advancement of periodontal disease. Furthermore, there is evidence to suggest that obesity may contribute to the acquisition and persistence of specific oral bacteria. Various studies have established a link between obesity, increased body weight, or higher body mass index (BMI), and the occurrence of periodontal diseases in humans. In a 2018 update to the classification of periodontal disease jointly conducted by the American Academy of Periodontology (AAP) and the European Federation of Periodontology (EFP), diabetes mellitus (DM) was included as a contributing factor that increases the risk of advancement in periodontal disease. [2, 3]. Certainly, obesity plays a crucial role in the development of type 2 diabetes (T2DM), which accounts for 90% of diabetes cases globally. The incidence of obesity is steadily increasing on a global scale. In 2016, 39% of adults were categorized as overweight (with a body mass index [BMI] ranging from 25 kg/m2 to less than 30 kg/m2), while 13% were classified as obese (with a BMI of 30 kg/m2 or higher). [4]. Throughout evolution, both animals and humans have developed a remarkable biological mechanism of storing excess energy in the form of fat to sustain life during periods of food scarcity. However, the modern lifestyle characterized by excessive calorie consumption and insufficient physical activity has led to an epidemic of obesity, which poses significant health risks. Epidemiological investigations have offered indications that obesity might be a potential risk factor in the onset of periodontal disease. [5]. Periodontitis has been proposed as a potential factor in the development of systemic conditions such as cardiovascular disease. This is primarily attributed to the inflammatory and microbial challenges posed by the inflamed surface of the periodontal pocket. [6]. Nesse and colleagues introduced a measurable framework for assessing the impact of chronic periodontitis, known as the "Periodontal inflammatory surface area (PISA)[7]. "The Periodontal Inflamed Surface Area (PISA) serves as a measure to quantify the square millimeter extent of the bleeding pocket epithelium. This assessment is derived from conventional clinical markers of periodontal health, specifically employing a combination of Bleeding on Probing (BOP) and either Periodontal Pocket Depth (PPD) or Clinical Attachment Loss and Gingival Recession.[8]. PISA functions as a valuable instrument for evaluating the extent and intensity of inflammatory lesions associated with periodontitis. Presently, several clinical studies are investigating the correlation between PISA and the susceptibility to systemic diseases.[9]. Furthermore, a connection exists between PISA and high-sensitivity C-reactive protein (hs-CRP), a marker indicating systemic inflammation. [10]. The objective of the research was to examine the connection





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between periodontitis, inflammatory scores, and obesity levels, as well as to quantify the inflammatory burden linked to chronic periodontitis by calculating the periodontal inflammatory surface area (PISA).

METHODOLOGY

Study Design

Individuals recruited for the study within the Department of Periodontics at Sree Balaji Dental College & Hospitalin Chennai, Tamil Nadu who fell into the categories of obesity or overweight participated. The study involved a survey based on questionnaires to assess aspects such as diet, lifestyle, and stress. Additionally, the height and weight of each participant were measured to calculate their Body Mass Index (BMI), with values exceeding 30.0 kg/m² indicating obesity[11]. Furthermore, waist-hip ratios were determined using measuring tape. The study also involved the measurement of various periodontal parameters, including bleeding on probing, clinical attachment level, recession, and pocket probing depth, which were assessed using a Williams probe. Moreover, specialized software analytic tools available at www.parsprototo.info were employed to deduce Periodontal Epithelial Surface Area (PESA) and Periodontal Inflammatory Surface Area (PISA).

Ethical Approval Number

The research underwent evaluation and received approval from the institutional ethics committee under the reference number (Sree Balaji Dental College & Hospital in Chennai, Tamil Nadu/IEC/12/2020/07) for our investigation. All participants were well-informed about the survey, and they willingly agreed to furnish the necessary data, providing informed consent prior to their involvement in the study.

Inclusion Criteria

Individuals between the ages of 25 and 60, of both genders, with a BMI greater than 25, and who have at least 20 scorable teeth were included in the study.

Exclusion Criteria

The study excludes individuals who fall into the following categories: pregnant and lactating mothers, those who have undergone antibiotic or steroid therapy within the last six weeks, individuals who have received periodontal treatment in the past six months, and those who are currently taking anti-depressants or have hormone imbalances.

Statistical Analysis

To ascertain how risk variables impacted the frequency of gum infection, a multivariate logistic regression analysis was conducted and the mean and SD of separate variables were assessed. At P 0.05, it was determined to be financially meaningful.

In four phases, the Periodontal Inflamed Surface Area was determined:

- 1. Based on six measurements taken around each tooth, a computer determined the mean pocket probing depth (PPD) for each tooth.
- 2. For that tooth, the mean PPD value was used to calculate the Periodontal Epithelial Surface Area (PESA).
- 3. PESA is determined by deducting Attachment Level Surface Area from Recession surface area (RSA).
- 4. The Periodontal Inflammatory Surface Area (PISA) value was determined by dividing the maximum number of sites by the product of the Periodontal Inflammatory Surface Area (PESA) of each tooth and the count of impacted sites exhibiting Bleeding on Probing (BOP).

Periodontal Inflammatory Surface Area of Individual Tooth PISA = (PESA × Number of sites with BOP) ÷ Maximum number of sites

PESA = Periodontal Epithelial Surface Area (surface area covered by the pocket epithelium) for the specific tooth.





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Number of sites with BOP = The number of sites around the tooth with Bleeding on Probing.

Maximum number of sites = The total number of sites that were probed around the tooth (usually six sites per tooth).

RESULTS

The main goal was to investigate the connections among periodontitis, inflammatory score, and obesity. The research included 21 participants and evaluated multiple factors such as diet, lifestyle, stress, BMI, and waist-hip ratio. Periodontal attributes, such as bleeding on probing, recession, and clinical attachment, were employed to determine Attachment Level Surface Area and Recession Surface Area. PESA and PISA were calculated using software tools. Given the limited sample size (21 participants), non-parametric approaches were employed for data analysis.

Descriptive

The aim is to investigate the correlation between age and PISA score.

Spearman's correlation was used because the assumptions of Pearson correlation were not met.

Interpretation

As the p-value exceeds 0.05, we can infer that there is no statistically significant association between age and PISA score.

The objective is to examine the correlation between BMI and PISA score.

Spearman correlation was used due to the violation of Pearson correlation's assumptions.

Interpretation

The fact that the p value is higher than 0.05 leads us to believe that there is no connection between BMI and PISA score.

The aim is to investigate the correlation between waist measurement and PISA score.

Spearman correlation was employed because the assumptions of Pearson correlation were not met.

Interpretation

As the p-value is greater than 0.05, we can infer that there is no statistically significant association between waist measurement and PISA score.

The objective was to examine the correlation between BOP and PISA score.

Spearman correlation was used due to the violation of Pearson correlation's assumptions.

Interpretation

As the p-value exceeds 0.05, we can infer that a noteworthy correlation between BOP and PISA score is lacking.

DISCUSSION

In recent times, clinical investigations have been conducted utilizing data related to the Periodontal Inflamed Surface Area (PISA), a metric used to quantify the extent of inflamed periodontal tissue. PISA measures the surface area of the epithelium in bleeding pockets [12]. Considering the documented association between PISA and glycated hemoglobin (HbA1c) [13,14], it serves as a valuable indicator for gauging the systemic impact of localized periodontal inflammation. Notably, recent findings have indicated a positive association between PISA and BMI (Body Mass Index) [15]. Conversely, Takeda and colleagues reported that in individuals with Type 2 Diabetes, PISA showed no significant correlation with obesity-related parameters [16]. Nonetheless, it is crucial to form a more extensive control cohort for a thorough examination of the association between PISA and obesity. Furthermore, multiple researchers have noted that although obesity doesn't stand alone as a risk factor for the advancement of periodontitis, there exist several common risk factors in the progression of both obesity and periodontitis. [17]. Currently, there is a scarcity of clinical evidence concerning the link between periodontal disease and overweight/obesity. Obesity, which is characterised by an excessive buildup of fat, is on the rise and is linked to





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chronic conditions including periodontitis [18]. The term "Infectobesity" refers to how alterations in microbiota brought on by persistent infection exacerbate circumstances in obese people. When it comes to the connection between periodontitis, obesity, and chronic illnesses, pro-inflammatory cytokines are a key player[19]. Gingivitis, corpulence, and persistent diseases are all closely related, and pro-inflammatory cytokines may be a major factor in this link[20]. The metabolic and immunological effects of obesity lead to an inflammatory state that makes people more vulnerable to periodontal disease[21]. Periodontal Inflammatory Surface Area, a novel metric, is used in this study to assess periodontal inflammation[22]. PISA estimates the inflammatory load and makes predictions about the likelihood that periodontitis may either cause or exacerbate other illnesses. Animal studies suggest probable processes including metabolic activity, hunger, and energy metabolism, and recent research have connected oral bacteria to obesity. Even a little increase in calorie intake without diet or exercise alterations may result in weight gain[23]. As a result, assessments of food intake and activity should be included in research relating to infection-associated weight gain. By measuring the inflammatory load brought on by periodontitis, ISA acts as a predictor for the risk that periodontitis may develop or exacerbate other illnesses[24]. The study relies on the pivotal variable in this research, the PISA index, established by Nesse.

Highlights of the Study

By assessing the inflamed surface area inside the periodontal tissues, this index serves as an indirect clinical diagnostic tool for detecting the inflammatory burden in chronic periodontitis. It is the first research to use the PISA approach to evaluate how obesity affects inflammation. The study found a link between increased BOP, which is seen as a sign of inflammation, and higher PISA scores.

Limitations of this Study

The primary limitations of this study include its small sample size, which may not accurately reflect all individuals, and its failure to show a significant correlation among adiposity and PISA score. Furthermore, it was impossible to properly account for confounding variables like mellitus and nicotine.

CONCLUSION

Future research in the field of Periodontal Inflammatory Surface Area (PISA) holds significant promise in advancing our understanding of periodontal disease and improving patient outcomes. One critical avenue for investigation is the potential of PISA scores as predictive indicators for disease progression. Examining the correlation between PISA and disease severity may offer valuable insights, enabling the development of more targeted and effective treatment strategies. Longitudinal studies are essential to explore how PISA scores correlate with treatment outcomes, providing a comprehensive understanding of the effectiveness of various therapeutic interventions in reducing inflammatory burden over time. Additionally, there is a need to explore the association between high PISA scores and systemic diseases such as cardiovascular disease, diabetes, or rheumatoid arthritis. This research could unveil the systemic implications of periodontal inflammation, guiding clinicians in considering broader health outcomes. Evaluating the feasibility and benefits of incorporating PISA measurements into routine clinical practice is pivotal. Such an integration could influence treatment planning and monitoring, addressing the need for standardized, cost-effective, and user-friendly methods. Furthermore, understanding the relationship between PISA and patient-reported outcomes is crucial, offering insights into the subjective experiences of individuals with periodontitis and enhancing holistic patient care. In essence, future research in these areas could revolutionize the diagnosis, treatment, and overall management of periodontal disease.

Authors Contribution Statement

Dr. V. Ramya led the entire research, which received approval from the institutional ethical committee at Sree Balaji Dental College. All the authors thoroughly reviewed and endorsed the final draft of the manuscript.





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Conflict of Interest

Conflict of interest declared none.

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[Table/Fig-1]: Descriptive Statistics								
Measurement	Sample Size	Minimum	Maximum	Mean	Standard Deviation			
BMI	21	25.10	35.80	28.2762	3.32128			
Age	21	24	57	40.57	8.829			
Waist (inches)	21	32.00	48.00	40.2762	4.42288			
PISA	21	8.944	1132.284	324.89476	268.046552			

[Table/Fig-2]: The correlation between age and PISA score.								
Correlations								
	PISA AGE							
	PISA AGE	Correlation coefficient	1.000	.288				
		Sig. (2 tailed)		.205				
Spearmans		N	21	21				
rho		Correlation coefficient	.288	1.000				
		Sig. (2 tailed)	.205	•				
		N	21	21				

[Table/Fig-3]	[Table/Fig-3]: The correlation between BMI and PISA score.								
Correlations	Correlations								
	BMI PISA								
		Correlation coefficient	1.000	327					
	BMI	Sig. (2 tailed)	•	.148					
Spearmans		N	21	21					
rho		Correlation coefficient	327	1.000					





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PISA	Sig. (2 tailed)	.148	
	N	21	21

[Table/Fig-4]: The correlation between waist measurement and PISA score.									
Correlations									
PISA WAIST									
		Correlation coefficient	1.000	.111					
	PISA	Sig. (2 tailed)	•	.632					
Spearmans		N	21	21					
rho		Correlation coefficient	.111	1.000					
	WAIST	Sig. (2 tailed)	.632						
	(inches)	N	21	21					

[Table/Fig-5]: The correlation between BOP and PISA score.								
Correlations								
	BOP PISA							
	BOP PISA	Correlation coefficient	1.000	.005				
		Sig. (2 tailed)		.982				
Spearmans		N	21	21				
rho		Correlation coefficient	.005	1.000				
		Sig. (2 tailed)	.982					
		N	21	21				

[Table/Fig-7	[Table/Fig-7]:Number of people who are obese by age, waist circumference, and body mass index.									
SCORE	1	2	3	4	5	6	7	8	9	10
PISA	8 to	120 to	232 to	354 to	466 to	578 to	690 to	802 to	914 to	1026 to
SCORE	120	232	354	466	578	690	802	914	1026	1138
BOP	15 to	24 to 32	32 to 41	41 to 50	50 to 59	59 to 68	68 to 77	77 to 86	86 to 95	95 to 104
	24									
WAIST	32 to	36 to 40	40 to 44	44 to 48						
	36									
BMI	25 to	30 to	35 to							
	29.9	34.9	39.9							
AGE	24 to	30 to 36	36 to 42	42 to 48	48 to 54	54 to 60				
	30									





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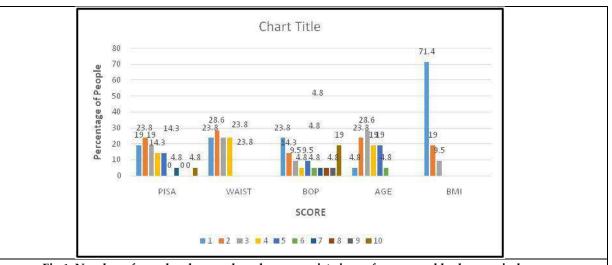


Fig 1: Number of people who are obese by age, waist circumference, and body mass index.

Hypothesis I: Oral bacteria increase metabolic efficiency Oral bacteria Food . GI Tract Increased fat increased storage metabolic efficiency Hypothesis II: Oral bacteria increase appetite Oral bacteria Increased Leptin and/or food Ghrelin consumption increased appetite Hypothesis III: Oral bacteria redirect energy metabolism TNF alpha → Increased Oral bacteria Insulin Adipocytes = blood fat glucose storage

Fig 2: Hypotheses about the role of oral bacteria in obesity.

Muscle -

Adiponectin





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RESEARCH ARTICLE

The Significance of Hormones in Governing the Physiology of **Menstrual Cycle**

Shital C. Shah*

Head, Department of Pharmacy, Ahmedabad Homeopathic Medical College, Parul University, Ahmedabad, Gujarat, India.

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*Address for Correspondence

Shital C. Shah

Head, Department of Pharmacy, Ahmedabad Homeopathic Medical College, Parul University, Ahmedabad, Gujarat, India.

Email: scshah1212@gmail.com



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ABSTRACT

The ovarian cycle is a critical component of female reproductive physiology, involving a series of hormonal and physical changes that prepare the ovaries for ovulation and fertilization. The ovarian cycle consists of two main phases: the follicular phase and the luteal phase. During the follicular phase, follicle-stimulating hormone (FSH), secreted by the anterior pituitary gland, stimulates the growth of ovarian follicles, each containing an immature egg (oocyte). As the follicles mature, they produce increasing levels of estrogen, which contributes to the proliferation of the uterine lining (endometrium) in preparation for implantation. Ovulation marks the transition between the follicular and luteal phases, triggered by a surge in luteinizing hormone (LH), also released by the anterior pituitary. The LH surge causes the dominant follicle to rupture, releasing the mature egg into the fallopian tube. The ruptured follicle then transforms into the corpus luteum, marking the start of the luteal phase. During the luteal phase, the corpus luteum secretes progesterone, which further enhances the uterine lining and maintains it for implantation. Progesterone levels remain high unless fertilization occurs. If fertilization does not take place, the corpus luteum degenerates, leading to a decline in progesterone and estrogen levels. This hormonal drop triggers the menstrual phase, wherein the uterine lining is shed, signaling the start of a new ovarian cycle. Hormonal regulation plays a crucial role in maintaining the ovarian cycle's rhythm and coordination. Gonadotropin-releasing hormone (GnRH) from the hypothalamus controls the release of FSH and LH, while estrogen and progesterone exert feedback control on the hypothalamus and pituitary to ensure proper cycle progression. Understanding the ovarian cycle's physiology is essential for comprehending female reproductive health and addressing conditions related to hormonal imbalances, ovulation disorders, and infertility.

Keywords: Gonadotropin, luteal phase, reproductive, physiology, fertilization





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INTRODUCTION

Ovary

The ovary are the paired oval sex glands with a length of 2-5 cm, a width of 1.5-3 cm, and a thickness of 0.5-1.5 cm, weighing about 10-20 g (average:14 g).

The ovary fulfils two major objectives:

- 1. generation of a fertilizable ovum
- 2. preparation of the endometrium for implantation through the sequential secretion of estradiol and progesterone

STRUCTURE OF OVARY

The ovary consists of three structurally distinct regions: an outer cortex containing the surface germinal epithelium and the follicles; a central medulla consisting of stroma; and a hilum around the area of attachment of the ovary to the mesovarium. The hilum is the point of attachment of the ovary to the mesovarium which contains nerves, blood vessels, and hilus cells having the potential to become active in steroidogenesis or to form androgen-secreting tumors. The outermost portion of the cortex, called the tunica albuginea, is covered by a single layer of surface cuboidal epithelium called the germinal epithelium.

The oocytes, enclosed in follicles are in the inner part of the cortex, embedded in stromal tissue. The stromal tissue is composed of connective tissue and interstitial cells, which are derived from mesenchymal cells have the ability to respond to LH with the production of androstenedione.

Granulosa Cell Layer

- 1. Membrane [Mural] Granulosa Cells: It forms the outermost layer of the membrana granulosa. The mural cells contain high levels of LH receptors and steroidogenic enzymes and account for most of the steroidogenesis in the follicle. Antral granulosa cells of membrane granulosa seem to be steroid genically and metabolically less active than mural granulosa cells.
- 2. Cumulus oophorus: It consist of the egg and a surrounding mass of granulosa cells. At ovulation, the granulosa cells of the cumulus oophorus are extruded with the egg, whereas the membrana granulosa becomes incorporated into the corpus luteum. The cumulus granulosa cells lack steroidogenic activity and have less LH receptors compared with the mural granulosa cells.

Theca interna layer, which forms just outside the basement membrane, shows typical steroidogenic features. They are responsible for producing the C19 (androgens) steroids that serves as a substrate for estrogen production. Theca externa is the outermost layer of the follicle, composed of fibroblasts, smooth muscle–like cells, and macrophages. It has important function during ovulation.

- 1. Primary interstitial cells: They constitute of C 19 -steroid producing cells located in the medullary compartment of the foetal ovary.
- 2. Theca-interstitial cells. Theca-interstitial cells represent the mature C 19 -steroid producing components of the follicle. It consists of theca interna and theca externa cells.
- 3. Secondary interstitial cells. Secondary interstitial cells are hypertrophied theca interna remnants surviving follicular atresia. These cells remain functionally and structurally unchanged.
- 4. Hilar interstitial cells: They are constituents of the ovarian hilum. They are large steroidogenic lutein-like cells. The secretory function of these cells is strongly supported by their prominence at the time of puberty, during pregnancy, and around menopause.

Two cell, two-gonadotropin theory of ovarian steroidogenesis:

This theory establishes that ovarian steroids are synthesized from cholesterol through the cooperative interactions of theca and granulosa cells. Theca cells: luteinizing hormone (LH) binds to luteinizing/chorionic gonadotropin receptor (LH/CGR) on the cell surface and stimulates the expression of the steroidogenic enzymes necessary for androgen production. Cholesterol is mobilized into mitochondria by steroidogenic acute regulatory protein (STAR)





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where it is converted to pregnenolone by cholesterol sidechain cleavage enzyme (CYP11A1). Pregnenolone diffuses into the smooth endoplasmic reticulum and is converted to progesterone by 3β -hydroxysteroid dehydrogenase (HSD3B). Progesterone is then converted to androstenedione by 17α -hydroxylase/17,20desmolase (CYP17A1). Granulosa cells: follicle-stimulating hormone (FSH) via signalling through follicle-stimulating hormone receptor (FSHR) stimulates the expression of enzymes necessary for estrogen synthesis. Androstenedione produced by theca cells diffuses into granulosa cells and is converted to testosterone by the enzyme 17β hydroxysteroid dehydrogenase (HSD17B) or to estrone by aromatase (CYP19A1). CYP19A1 utilizes testosterone to produce 17β -estradiol. However, HSD17B can also produce 17β -estradiol using estrone as a substrate. The classical two cell theory is supported by molecular finding, ovarian steroidogeneses in the pre ovulatory follicle takes place through LH receptor on theca and FSH receptor on granulosa cells cyclic amp production and increase SF1 binding to multiple steroidogenic promoter mediate LH action in theca cells, the pre ovulatory follicle produces estradiol through parakine interaction between theca and granulosa cell in response to stimulating steroidogenic factor 1 (sf1) a member of the orphan nuclear receptor family x as a master which to initiate transcription of seriace of steroidogenic activity of each cell type.

Meiotic and mitosis

The first meiotic division in female ovary occurs before the birth of female child during the gonadal development of foetus in intrauterine life. The primordial germ cells are known to migrate from the yolk sac to the genital ridge by amoeboid movements with the aid of pseudopodia. By the fifth week of gestation it arrives at the genital ridge, known as oogonia. At about 8 weeks of intrauterine life, persistent mitosis increases the total number of oogonia to 600,000. From this point on, the oogonial development is subject to three simultaneous processes: mitosis, meiosis, and oogonial atresia. Between weeks 8 and 13 of fetal life, some of the oogonia depart from the mitotic cycle and enter the prophase of the first meiotic division.

The number of primary oocyte is maximum at 20 weeks of gestation that means at 5 months of intra uterine life, and that is about 6-7 million.

At birth = 1 to 2 million

At puberty= 3 to 5 lakhs

Out of this only 500 primary oocyte will be matured during the reproductive age of female under the influence of Gonadotropins-FSH & LH and ovarian steroids. Every month around 15 to 20 primordial follicles convert into primary follicles. In the beginning of the normal menstrual cycle there is a high FSH. It helps to start the growth of the primordial follicles into primary follicles and under the influence this FSH level, it allows only one primary follicle to mature into secondary follicle (Antral follicle). This follicle will grow bigger and bigger and will mature as graafian follicle.

DEVELOPMENTAL STAGES OF THE OVARIAN FOLLICLE:

PRIMORDIAL FOLLICLES are located in the periphery of the cortex and inferior to tunica albuginea, surrounded by a single layer of squamous follicular cell containing immature primary oocytes. Every month around 15 to 20 primordial follicles convert into primary follicles under the effect of FSH. These cells give nourishment to ovum

PRIMARY OOCYTE: Primordial follicles start to grow at puberty under the influence of FSH. The outermost layer of follicular cells rests on a well defined basement membrane that separates it from the ovarian stroma. The oocyte is separated from the surrounding follicular cells by a glycoprotein layer called zona pellucida. The connective tissue stroma surrounding the follicle begins to condense and form theca follicular.

SECONDARY OOCYTE: secondary oocye also known as antral follicle. Secondary oocyte is surrounded by zona pellucida. Cuboidal granulosa cell layer in secondary follicle secretes estradiol. Irregular fluid filled space among the follicular cells join to form antrum. Theca follicula later differentiate into 2 layers in the secondary follicle. They are theca externa and theca interna. Follicular cells and theca interna secretes estradiol.





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TERTIARY FOLLICLE: The follicle increases in size.

GRAFFIAN FOLLICLE: The mature Graafian follicle is spheroidal or ovoid in shape and seen bulging out of the cortex in surface of the ovary, contains pent-up secretion, the liquor folliculi. The granulosa cells are collected together to form a projection into the cavity of the Graafian follicle. This projection is called as Cumulus oophorus (discus proligerus). The ovum lies within cumulus oophprus is known as secondary oocyte. The granulosa cells layer surrounded around the ovum is known as corona radiata. The theca interna cells enlarge during the maturation of the follicle and shortly before ovulation they are larger than the granulosa cells. The graafian follicle grows at the rate of 1-2mm daily and attains the size of 20mm or more at ovulation.

HORMONAL MECHANISM: OVARY

LH AND FSH SYNTHESIS:

Gonadotroph cells of pituitary gland synthesis LH and FSH, they are made up of 2 peptide subunits called Alfa and beta, each subunit contains cysteine, multiple disulphide linkages, multiple carbohydrate -molecules and help in the metabolism of these hormones, on binding of GNRH to its receptor the biosynthesis of the gonadotropins proceeds by transcriptions of the subunit genes, mRNAs- post translational modification of the precursor subunits, mature hormone packaging and secretion. The alpha subunit of LH and FSH have polypeptide structure and beta subunit have amino acid structure

FUNCTION of FSH- FSH receptors have localized into granulosa cells and stimulate granulosa cells to synthesis aromatase which converts androgens (androstenedione and DHEA) to estradiol. Necessary for normal follicular development, ovulation, fertility and estrogen production, during the follicular phase of the menstrual cycle, a mature dominant follicle secrets estradiol and inhibin, send a negative feedback to hypothalamus for suppressed FSH secretion. FSH remains low throughout the luteal phase preventing the development of the new follicle. The granulosa cell LH receptor requires the continued presence of FSH for its function.

Luteinizing function of ovary:

LH is essential for complete follicular development, formation of corpus luteum and production of androgens, it plays a major role in the promotion of the theca interstitial cell, increase LH receptor and key substance for steroidogenic process . The preovulatory follicle secrets estradiol during the follicular phase of the menstrual cycle, the estrogen synthesis in the ovary by the aromatization process. Ovarian steroidogenesis (estrogen and progesterone) in the preovulatory follicle takes place through LH receptor on theca cells and FSH receptor on granulosa cells. The star protein is the primary regulator of production of androstenedione, which diffuses into granulosa cells as a precursor for the estrogen, in the preovulatory follicle cholesterol in a theca cells arises from circulating lipoprotein.

LH and FSH RECEPTORS

Peptide hormone Receptors are large proteins whose essential functions are:

- a) the ability to bind the hormone
- b) the ability to couple hormone binding to hormone action.

Some receptors are located within the cell, and function as transcription factors (steroid hormones). Other receptors are located on the cell surface and function primarily to transport their ligands into the cell by a process referred to as receptor mediated endocytosis. The cell surface receptor triggers intracellular signaling pathways.

Ligands are the substances which binds with receptor. E.g. all hormones acting as a ligand in the body. The point where the ligand binds with the receptor are called 'ligand binding domain'. Once ligand binds with the receptor, is supposed to produce some biological action.





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G-protein-coupled receptors (GPCRs):

G-protein or guanosine nucleotide binding proteins act as exchange factors. In terms of structure, GPCRs are characterized by an extracellular N-terminus, followed by seven transmembrane (7-TM) α -helices (TM-1 to TM-7) connected by three intracellular (IL-1 to IL-3) and three extracellular loops (EL-1 to EL-3), and finally an intracellular C-terminus. The GPCR arranges itself into a tertiary structure resembling a barrel, with the seven transmembrane helices forming a cavity. They act as cell surface receptors that detect molecules outside the cell and activate cellular responses. They are coupled with G proteins hence, they pass through the cell membrane forming six loops out of which 3 are extracellular loops and 3 intracellular loops. Ligands can either bind to the extracellular N-terminus and loops (e.g. glutamate receptors) or to the binding site within transmembrane helices. They are mostly activated by agonists, although a spontaneous auto-activation of an empty receptor has also been observed.

There are principally two signal transduction pathways involving the G protein-coupled receptors:

- the cAMP signal pathway
- the phosphatidylinositol signal pathway.

A conformational change brought about by ligand binding to the GPCR enables the GPCR to function as a guanine nucleotide exchange factor (GEF). The GDP bound to the G protein is then exchanged for a GTP by the GPCR, activating the corresponding G protein. Depending on the type of α subunit, the G protein's α subunit can target functional proteins directly or further affect intracellular signaling proteins by dissociating from the β and γ subunits along with the bound GTP. FSH and GnRH ligands are binded to one or more receptors on GPCRs.

ESTROGEN SYNTHESIS

The naturally occurring estrogens are C18 steroids characterized by the presence of an aromatic ring, a phenolic hydroxyl group at C3, and a hydroxyl group (estradiol) or a ketone group (estrone) at C17.

Estrogen is produced in at least three major sites:

- (1) by direct secretion from the ovary in reproductive-age women;
- (2) by conversion of circulating androstenedione, originating from the adrenal or ovary or both, to estrone in peripheral tissues; and
- (3) by conversion of androstenedione to Estrogen in estrogen target tissues.

In the latter two cases, the same tissue converts the weak estrone to estrogen. For estrogen to form at these sites, genes encoding the enzymes aromatase and reductive 17β -hydroxysteroid dehydrogenases (17β -HSD) must be expressed. Multiple genes with overlapping functions may produce protein products that contribute to reductive 17β -HSD activity in peripheral tissues. A unique reductive 17β -HSD enzyme called HSD17B1 is encoded by a particular gene that is mostly expressed in the ovary. There is only one gene that codes for aromatase (CYP19A1). It is especially crucial for postmenopausal women and those suffering from diseases that depend on estrogen, like endometriosis, breast cancer, or endometrial cancer, to produce estrogen through peripheral and local conversion.

Estrogen nuclear receptor

A family of ligand-regulated transcription factors known as nuclear receptors is triggered by a variety of lipid-soluble signals, such as thyroid hormone, retinoic acid, and oxysterols, as well as steroid hormones like progesterone and estrogen. In contrast to the majority of intercellular messengers, the ligands do not need to act through cell surface receptors in order to cross the plasma membrane and interact with nuclear receptors inside the cell. Upon activation, nuclear receptors exert direct control over the transcription of genes that govern an extensive range of biological functions, such as cell division, proliferation, metabolism, and reproduction. While their primary role is as transcription factors, some nuclear receptors have also been discovered to control cellular processes in the cytoplasm. For instance, endothelial cells cytoplasm contains an estrogen receptor, which estrogens use to quickly activate signaling pathways that regulate endothelial cell migration and vascular tone.

Phosphorylation is one of the main mechanisms regulating the activities of orphan receptors and can activate certain nuclear receptors independently of ligand binding. Phosphorylation is another posttranslational modification that





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can modify the functions of nuclear receptors. Given the wide variety of processes controlled by nuclear receptors, their dysregulation can contribute to numerous diseases, including cancer, diabetes, and infertility. However, because they bind to small molecules, they represent promising therapeutic targets for which selective agonists and antagonists can be engineered Because nuclear receptors regulate many genes in many tissues, synthetic ligands usually show beneficial therapeutic effects and unwanted side effects that limit clinical use. Major goals in the nuclear receptor field therefore include attaining a better understanding of the mechanisms underlying their actions in specific cell types and ways in which to selectively modulate their activities. A characteristic feature of nuclear receptors with respect to their integrative roles in development and homeostasis is their ability to regulate different genes in different cell types. For example, estrogen receptors regulate different sets of genes in the brain, breast, and uterus that contribute to the distinct functions of those organs. Recent studies indicate that tissue-specific responses are a consequence of binding of nuclear receptors to enhancer elements that are selected in a cell-specific manner.

OVARIAN CYCLE

Hypothalamic-pituitary-ovarian Axisand synchronisation of this Axis is necessary for ovulatory menstrual cycle in female reproductive age. Regular ovulation, predictable menses and other female reproductive functions are coordinated by the functions of the Hypothalamus, Pituitary gland, Ovaries and Endometrium. Arcuate nucleus and preoptic nucleus of hypothalamus have special neurons which release. The hypophyseal portal systemtransports GnRH from hypothalamus to anterior pituitary and stimulates gonadotroph cells. The pulsatile secretion of GnRHinfluences Gonadotroph cells of the pituitary gland & secrets gonadotropins i.e. FSH and LH. The periodicity and pulsatile rhythm of GnRH and gonadotropin secretion are necessary in regulating reproductive system. Cyclic activity in ovary is controlled by an interaction between estradiol, progesterone, FSH and LH.

On the 1st day of menstrual cycle, small and medium sized follicles are present in the ovaries, so low level of estradiol, GnRH pulse frequency is relatively fast(90 minutes). As estradiol level increases in the follicular phase, GnRH pulse frequency is relatively fast(60 minutes) in a mid-follicular phase. The Estradiol makes the endometrium to proliferate. Before the day of ovulation the level of the estradiol is at its maximum level. And send the negative feedback to the hypothalamus to stop the FSH secretion & at the same time a positive feedbackaction is send to the hypothalamus to trigger a LH surge. This LH surge influences ovulation.

The corpus luteum starts to secret large amount of Progesterone (40mg/day) and some estradiol during the midluteal phase of the ovarian cycle. High level of the progesterone decrease GnRH pulsatility(1 pulse in 100 minutes) and (1 pulse in 200 minutes) during late luteal phase. The functional lifespan of the corpus luteum is normally 14 days, thereafter the corpus luteum spontaneously regresses, unless pregnancy occurs. As the corpus luteum regresses, progesterone and estrogen secretion diminishes. Reduced progesterone sends the negative feedbacksignals to the hypothalamus, and allows secretion of FSH. Reduced progesterone is also a withdrawal support to the endometrium lining as a result the endometrium is shed and bleeding starts and a new cycle of the menses begins.

Definition

Ovulation consists of rapid follicular enlargement followed by outcoming of the follicle from the surface of the ovarian cortex. Rapid follicular enlargement and follicle extrusion from the ovarian cortex's surface characterize ovulation. The rupture of the follicle and the extrusion of an egg come next. It happens when the corona radiata-enclosed ovum breaks free from the graafian follicle. The sharp increase in the amount of estradiol in the blood causes ovulation. The midcycle surge in LH is caused by the hypothalamus receiving positive feedback from an increase in estradiol levels. Ovulation requires a peak LH concentration of 75 ng/ml. 34–36 hours after the beginning of the LH surge, ovulation happens predictably. There is antral fluid and the ovum explodes when there is a rupture. When ovulation occurs there is a sudden shrinkage in the size of a follicle and appearance of free fluid in the pouch of douglas. After ovulation graafian follicle collapses and gets transformed into temporary endocrine organ called as Corpus Luteum. During the first few hours after the explosion of the ovum from the follicle, the remaining granulosa cells and theca cells change rapidly into lutein cells. The lutein cells derived from the granulosa layer forms the





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granulosa lutein cells. The lutein cells derived from the theca layer forms the paralutein cells. These cells undergo process of hypertrophy and hyperplasia called luteinisation. The corpus luteum is the endocrine gland that serves as the major source of Progesterone and Estrogen secreted by the ovary during the luteal phase. corpus luteum proliferates for about 12 days. The granulosa lutein cells become spindle-shaped and secrets progesterone and estrogen hormones. The theca lutein cells form androgens- Androstenedione and Testosterone. Before rupture of the follicle the granulosa layer is not vascularised. Corpus luteum formation is the penetration of the follicle basement membrane by the blood vessels which provides the granulosa lutein cells with low density lipo-protein(LDL). LDL cholesterol serves as the substrate for corpus luteum progesterone production. A key factor of steroidogenesis in the corpus luteum is LH. LH-receptor is maintained throughout the functional life span of corpora lutea. The functional life span of the corpus luteum is 14 days. There after the corpus luteum spontaneously regresses. It is replaced, unless pregnancy occurs, by an avascular scar referred to as the corpus albicans. Progesterone levels will drop

SECRETORY PHASE

Progesterone and estrogen, both together are secreted in large quantity by the corpus luteum. The estrogen cause additional proliferation in the endometrium. The endometrium is divided into an upper two-third functionalis layer and a lower one-third basalis layer. The purpose of the functionalis layer is to prepare for the implantation of the blastocyst; therefore it is the site of proliferation, secretion and degeneration. The purpose of the basalis layer is to provide the regerative endometrium following mestrual loss of the functionalis layer. The endometrium swells and secretes more when progesterone is present. The glands become more twisted. An excess of secretory materials builds up within the glandular epithelial cells. Lipid and glycogen levels in the stromal cell's cytoplasm rise significantly, and the blood supply rises along with them. The spiral arteries and their capillaries pierce the entire thickness of the endometrium. Endometrial tissue is shed during menstruation along with steroid hormone hemorrhage. The corpus luteum in the ovary abruptly involutes (decreases in size after an enlargement) if the ovum is not fertilized, which occurs roughly two days before the end of the monthly cycle. Additionally, the levels of secretion of the ovarian hormones progesterone and estrogen drop

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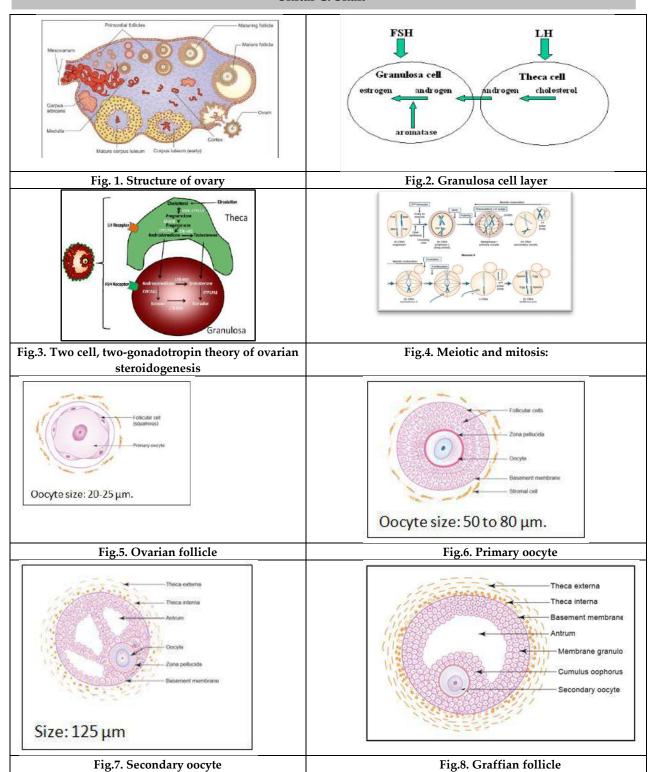
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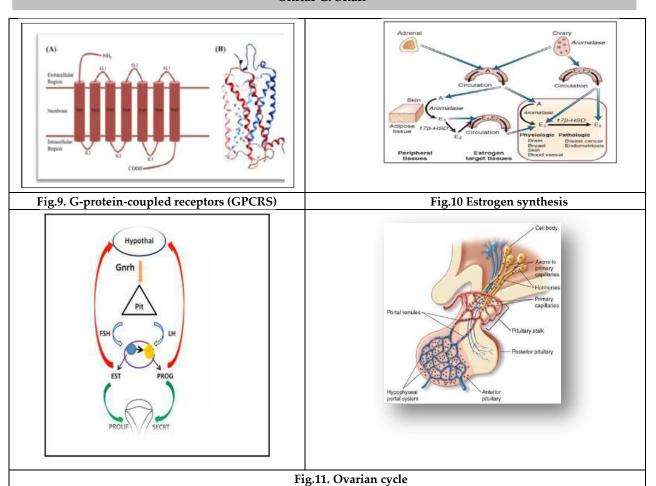
Shital C. Shah







Shital C. Shah







RESEARCH ARTICLE

Effect of Serratus Anterior Muscle Strengthening on Pain and Function External Shoulder Impingement Syndrome: Secondary **Experimental Study**

Priyal Ninama^{1*}, Rushi Gajjar² and Gaurav Patel³

¹Student, Department of Physiotherapy, Ahmedabad Physiotherapy College, Parul University, Gujarat,

²Associate Professor, Department of Physiotherapy, Ahmedabad Physiotherapy College, Parul University, Gujarat, India.

³Principal, Department of Physiotherapy, Ahmedabad Physiotherapy College, Parul University, Gujarat, India.

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*Address for Correspondence Priyal Ninama

Student, Department of Physiotherapy, Ahmedabad Physiotherapy College, Parul University, Gujarat, India.



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ABSTRACT

shoulder pain is one of the most common complaints of patient's world wide, especially in older age group. SIS is the syndrome associated with compression and inflammation of supraspinatus tendon as it passes through sub acromial space. There for to investigate the serratus anterior muscle strengthening exercise + scapular muscle strengthening exercise programe to improve balance of the shoulder and neck, posture, ADL'S, QOL is the main focus of this study. 32 subjects were participants in this study based on inclusion and exclusion criterias with the age of 20 to 50 year. Pre and Post SPADI and NPRS was checked. The serratus anterior muscle strengthening exercise+scapular muscle strengthening exercise, control group continued their scapular muscle strengthening exercises. The outcome measures taken at baseline and after 2 weeks of intervention. Intervention shows significant results in serratus anterior muscle exercise + scapular muscle strengthening exercise had significant effect on NPRS score and SPADI score in secondary external impingement syndrome in Group A. This study showed that there is a positive effect of Serratus Anterior muscle strengthening exercise and Scapular muscle strengthening exercise is more favorable when compared to scapular muscle strengthening exercise.

Keywords: Shoulder impingement syndrome, serratus anterior muscle, secondry external impengement, NPRS, SPADI





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INTRODUCTION

One of the most common musculoskeletal concerns is shoulder pain. With a prevalence of %44–65 among all shoulder problems, subacromial impingement syndrome (SIS), also known as shoulder impingement syndrome, is the most typical cause of shoulder pain.[1] The labrum also enhances stability by deepening the concavity of the glenoid socket. The capsule tightens or "winds up" in various extremes of position; for example, the inferior pouch tightens in extreme abduction and external rotation, serving to stabilize the joint.[2]The coracoacromial arch and humerus are what causes the rotator cuff and subacromial bursa to mechanically compress. Subacromial impingement, sometimes known as "external impingement," internal impingement, which can be further classified into anterior or posterior impingement.[3]Movement is painfully restricted, there is a functional impairment, and everyday activities are restricted. When the arm is elevated during unconstrained overhead reaching, flexion, abduction, or scapular plane abduction, the scapula will normally rotate upward and slant posteriorly on the thorax.[4] The subacromial impingement syndrome has both primary and secondary forms.[5] Secondary impingement happens when the humeral head's functional centering is disturbed, such as by muscle imbalance. This abnormal displacement of the centre of rotation in elevation causes soft tissue entrapmentA widely used classification system for acromial shape is flat (type I), curved (type III), or hooked (type III).[6]

SERRATUS ANTERIOR MUSCLE

The fan-shaped serratus anterior starts on the superolateral surfaces of the first to eighth or first to ninth ribs at the lateral wall of the thorax and inserts along the superior angle, medial border, and inferior angle of the scapula.[7] The scapula is pulled forward around the thorax by the serratus anterior muscle, allowing the arm to be extended and anteverted.[8] The serratus anterior starts to separate from the mesenchyme present at the ventral ends of the lower cervical myotomes in the 9 mm embryo.[9] The lower part of the serratus anterior and the latissimus dorsi are supplied by the thoracodorsal artery.[10] The innervation of the serratus anterior is carried out by the long thoracic nerve.[11] For the scapulohumeral rhythm to be maintained when raising the arm, the serratus anterior muscle must be activated appropriately. The serratus anterior helps the scapula rotate upward and tip posteriorly when the humerus is raised. It also keeps the scapula level on the thorax, preventing scapular winging. One mechanism behind shoulder impingement syndromes has been proposed to be impaired serratus anterior activation, which has been linked to altered scapular kinematics. [12] A self-report questionnaire for people with shoulder discomfort called the Shoulder Pain and Disability Index (SPADI) has 13 questions and is separated into two domains: disability (5 items) and pain (8 items). A higher score denotes poorer shoulder function and discomfort. For the SPADI, the ICC value is 0.89.[13]Unidimensional measurement of pain intensity is provided by the Numeric Pain Rating Scale (NPRS). The 11-point numeric scale runs from "0" (i.e., "no pain") to "10," which denotes the most intense agony. For the NPRS, the ICC value is 0.92.[14] Hence, the purpose of the present study is to evaluate the effect of serratus anterior muscle strengthening on pain and function in secondary external impingement syndrome.

MATERIAL AND METHODOLOGY

32 subjects were taken from sainath hospital bopal-ahmedabad by using simple random sampling for this experimental study of 2 weeks duration.

Inclusion criteria

Age between 20 to 50 years. Male and females both are included. Positive Neer's, Howkins-Kennedy, Yokum test and apprehension test. Patient willing to participate. Shoulder pain and disability is more than 40%.

Exclusion criteria

Any traumatic injury to bone such as fracture, dislocation. Herniation of cervical disc with or without radiating symptoms. Pain with shoulder movement under 60 degrees. Infection of bone or soft tissues such as abscess. Higher functions are affected in subject.





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Procedure of outcome measure

A self-report questionnaire for people with shoulder discomfort called the Shoulder Pain and Disability Index (SPADI) has 13 questions and is separated into two domains: disability (5 items) and pain (8 items). A higher score denotes poorer shoulder function and discomfort. For the SPADI, the ICC value is 0.89.[13] Unidimensional measurement of pain intensity is provided by the Numeric Pain Rating Scale (NPRS). The 11-point numeric scale runs from "0" (i.e., "no pain") to "10," which denotes the most intense agony. For the NPRS, the ICC value is 0.92.[14]

Procedure of intervention

Group A:(EXPERIMENTAL GROUP) SERRATUS ANTERIOR MUSCLE STRENGTHENING + SCAPULAR MUSCLE STRENGTHENING EXERCISE

This group received serratus anterior muscle strengthening exercise and scapular muscle strengthening exercise for 6 days per week for 2 week .3 sets of 10 repetitions between 1 minute rest up exercise.

- 1) serratus anterior muscle strengthening exercise
 - a) The wall slide
 - b) scapular plane shoulder elevation
- 2) scapular muscle strengthening exercise
 - a) side lying forward flexion
 - b) side lying external rotation
 - c) prone horizontal abduction with external rotation
 - d) prone extension in neutral position with weights

Group B: (CONTROL GROUP) SCAPULAR MUSCLE STRENGTHENING EXERCISE

- a) side lying forward flexion
- b) side lying external rotation
- c) prone horizontal abduction with external rotation
- d) prone extension in neutral position with weights

RESULT

Statistical analysis was performed using SPSS version 26. Total thirty two subjects are selected for two weeks of training protocol. In this study Group A, received Serratus anterior muscle strengthening exercise and scapular muscle strengthening exercise (n=16) and Group B received Scapular muscle strengthening exercise (n=16). Both the groups were similar in age, gender and function. The study analysis was done to evaluate between group interventions, Pre-post intervention. Within group, The effects were explored using paired t-test and Between group the effects were explored by using independent t-test. In the result data shows value of mean, standard deviation (SD), t value and degree of freedom (df). Statistical significance was set at p<0.001. From total 32 subjects, there were 7 females and 9 males in the study. They were equally divided in both group. By using independent sample test results for NPRS and SPADI shows significant improvement in GROUP A(serratus anterior muscle strengthening exercise and scapular muscle strengthening exercise) as its p value is <0.001. (Graph 1,2,3) (Table 1)

As shown in table 2 within group result by using paired t test shows significant results in NPRS and SPADI in both the groups except as its p value is <0.001.

DISCUSSION

The of impact serratus anterior muscle strengthening exercise and scapular muscle strengthening exercise versus scapular muscle strengthening exercise on pain and function in secondry external impingement syndrome. The pain was evaluated using the Numeric Pain Rating Scale (NPRS), Disability was assessed using Shoulder Pain and Disability Index questionnaire scale (SPADI) were used as a primary outcome, whereas Hawkins Kennedy Test, Positive Neer's Test, Yokum Test, Apprehension Test was used as a secondry outcome measure. A study by Dvir &





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Berme 1978, Johnson et al. 1994. The biggest moment arm for producing scapular upward rotation torque is found in the serratus anterior. The serratus anterior line of action can also make a significant contribution to scapular posterior tilting. Ekstrom Ra, et al. 2003 study concluded The serratus anterior is a muscle that is first activated above 90 degrees of shoulder flexion, although our study is not the first to look at this. The serratus anterior was most fully stimulated by raising the shoulder over 120 degrees in the plane of the scapula while utilising a hand weight. Mc Clure Pw, et al. 2004 study concluded When compared to the push-up plus exercise against the wall, the wall slide and scapular plane shoulder elevation exercise more fully engaged the upper trapezius. This might be due to the upper trapezius acting as an agonist for the wall slide and scapular plane shoulder elevation necessary for proper performance. Carrie M Hall, et al. 1980 the main stabilizer of the scapula include serratus anterior, all 3 trapezius muscles, rhomboids and levator scapulae. If there is weakness or imbalance of these muscle, it may lead to altered position of the scapula and may also decreased its stability. As a result of this, the function of the shoulder complex may become inefficient leading to decrease in neuromuscular perfomance which may have been corrected by improving the strength of the lower trapezius and serratus anterior muscle as found in our study , thereby decreasing the load on the cervical efficient muscle joint complex. Blackburn Ta, et al.1990 study concluded that Blackburn has described the prone full can, or horizontal abduction (100 degree of elevation) with external rotation, as an exercise that facilitates high supraspinatus electromyographic activity. Because the infraspinatus and teres minor have very similar concentric muscle actions of externally rotating the humerus they can generally be exercised with same movements.

CONCLUSION

This study showed that there is a positive effect of serratus anterior muscle strengthening and scapular muscle strengthening exercise programe to effects on pain reduction and improved function and increased rang of motion or strength in secondry external impingement syndrome but Group A showed greater significant difference as compared to Group B. Serratus Anterior muscle strengthening exercise and Scapular muscle strengthening exercise is more favorable when compared to scapular muscle strengthening exercise.

Clinical implication

This study shows positive effect of serratus anterior muscle strngthening exercise and scapular strengthening exercise programme include strngth, range of motion, balance, posture, quality of life, ADL's. so, Secondary external impingement subjects use this techniques are improved performance and to reduced the risk of injuries.

Limitation

Long term follow up was not taken up for this study.

Future recommendation

A large sample size would be recommended in future reserches. More subjects can be taken for study. Interpretations can be done for long periods. studies for the long term effect of this treatment could be concluded.

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Table 1: independent sample test for between group result

	Independent Sample test								
	Т	df	Sig.(2 tailed)	Mean difference					
A.D		38	0.002	-0.90000					
ABnprs	-3.421	36.570	0.002	-0.90000					
ABspadi	-3.919	38	0.000	-5.40000					
Tibopuui		38	0.000	-5.40000					

Table 2: Paired sample test of within group results

Table 2.1 affect sample test of within group festilis											
Paired sample test											
Group			Mean	Standard deviation	Т	Sig (two tailed)					
Group			wican	Startdard de viation	1	Sig (two tailed)					
	Pair 1	Anprs Pre	7.5500	1.05006	23.134	0.000					
	Pair 1	Anprs Post	2.3500	0.74516	23.134						
Group A	D. i. o	Aspadi pre	76.7500	15.02235	15.074	0.000					
	Pair 2	Aspadi post	20.1500	4.49883	15.874						
Group B	Pair 1	Bnprs Pre	7.2500	1.37171	15.292	0.000					

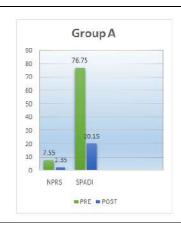




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	Bnprs Post	3.2500	0.91047		
Pair 2	Bspadi pre	74.7000	12.27792	17.914	0.000
1 all 2	Bspadi post	25.5500	4.21120	17.714	



Group B

70

70

60

50

40

30

25.55

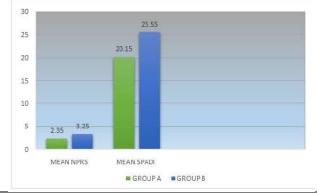
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NPRS SPADI

PRE POST

Graph 1: Group A Pre and Post Mean of NPRS and SPADI

Graph 2: Group B Pre and Post Mean of NPRS and SPADI





Graph 3: NPRS and SPADI Between Two Groups

Photograph 1: The Wall Slide & Scapular Plane Shoulder Elevation



Photograph 2: 1)Prone Horizontal Abduction With External Rotation 2) Side Lying Forward Flexion 3)Side Lying External Rotation 4) Prone Extension In Neutral Position With Weights





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RESEARCH ARTICLE

Physico -Chemical assessment of Groundwater quality in Bagepalli taluk of Chikballapur district, Karnataka

Ramesha Iyyanahalli^{1*} and Kiran B.R.²

^{1*}Associate Professor, Department of Zoology, Government Science College, Chitradurga-577501, India. ²Research & Teaching Assistant in Environmental Science, DDE Kuvempu University, Shankaraghatta-577451, India.

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*Address for Correspondence Ramesha Iyyanahalli

Associate Professor, Department of Zoology, Government Science College, Chitradurga-577501, India.



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ABSTRACT

The present observation deals with the assessment of groundwater quality in Bagepalli taluk of Chikballapur district, Karnataka, India. To analyze the physico-chemical characteristics groundwater samples and assess the suitability of water for human consumption and irrigation. Also to find out the source of pollution. Methods: A total of 10 water samples (13.78°N latitude and 77.79°E longitude) (S1 to S10) were collected from selected village borewells of Bagepalli taluk, Chikkaballapur District during June -2010 to May-2011. The temperature as recorded at the time of sampling by using mercury thermometer. The pH was measured with pH meter. All the other parameters were measured as per the Standard Methods (APHA-2005). One-way ANOVA with post-Hoc Tukey HSD tests were carried out by using asatsa.com software. The water temperature fluctuated between 25 O C and 35 O C. pH values varied from 7.0 to 8.8. The values of electrical conductivity ranged 302 µmhos/cm to 2718 µmhos/cm. TDS values deviated from 311 mg/L to 2483 mg/L Total hardness varied from 311 mg/L to 1042 mg/. While, alkalinity values ranged from 200 mg/L to 588 mg/and chloride values from 80 mg/L to 527 mg/L Nitrate data fluctuated from 48 mg/L to 270 mg/L. Sulphate values deviated from 10 mg/L to 95 mg/L. However, phosphate varied from 0.2mg/L to 1.4 mg/L. Sodium, potassium and iron levels fluctuated from 70-360 mg/L, 1.2-4.2 mg/l and 0.1-0.7 mg/l respectively. The groundwater of Bagepalli taluk had varied in different seasons due to anthropogenic activities, deterioration by agricultural activities, over exploitation and also of drought circumstance. The consequences of all the above findings are discussed by emphasizing for suitable water quality management. These results contribute to environmental decision making in the management of ground water resources. In most of the ground water samples the values of electrical conductivity, total hardness, nitrate, total alkalinity, and chloride are exceeded the permissible limits of BIS





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standards. It is due to seepage of sewage and urban runoff into the ground waters. Other factors include lack of proper sewerage facilities and open drains in the study area. Hence, proper treatment is necessary before utilized for domestic purposes..It is significant to carry out measures for augmenting groundwater availability for use and increasing use effectiveness, the aspect of improving groundwater governance. This current study contributes to the assessment of ground water quality by policy makers and stakeholders, enabling the arrangement and managing the ground water resources in Bagepalli taluk of Chikaballapura district.

Keywords: Ground water quality, Bagepalli taluk, Chikballapur, BIS standards, Permissible limits, Karnataka.

INTRODUCTION

Ground water is the major resource of drinking water in both urban and rural India. Discovering dimensions of ground water quality is an important role for the past two decades because of the uncertainties of water resources, expansion of pollution level through the population growth and industrial development. On the other hand, ground water system receives the soluble inorganic and organic pollutants through the percolation of vast domestic sewage and industrial effluents. Various hydro-geochemical process determines diverse physico-chemical characteristics of ground water chiefly in the region of saturation (Islam et al.,2017a; Ahmed et al.,2020;Bhuiyan et al.,2016). Ground Water is an essential natural supply of fresh water worldwide. Due to the water's suitability for a variety of uses, including drinking, livestock, irrigation, aquaculture, mining, and industrial uses, groundwater quality is just as important as quantity. Lately, emerging nations like India confronting the interest of unadulterated water for drinking and different designs are expanded quickly because of absence of ground water accessibility in the greater part of the areas. The central point that influences the nature of water because of quick increment of industrialization and rural improvement exercises and urbanization, which causes ground water gets debased (Juahir et al.,2011).

According to the WHO (1985), approximately 80% of human diseases are transmitted through water. In arid and semiarid regions, where surface water is scarce and unevenly distributed, approximately one third of the freshwater used to meet all domestic, industrial, and agricultural needs comes from groundwater (Liang Chen et al.,2017). Water quality assessment is usually cited as a limiting factor in proposing certain anthropogenic activities in a region (Jose Galizia et al., 2016; Fenge Zhang et al., 2019; Daniele Parrone et al., 2020). As a result, understanding the physical, chemical, and biological properties of groundwater resources is crucial to determining their applications: human utilization, water system, modern purposes, and others (Fenge Zhang et al., 2019; Daniele Parrone et al., 2020; Franklin et al., 2019; Silva et al., 2021) Various earlier studies were carried on assessment of ground water quality and pollution in different parts of the India and the world. They come across significant deterioration in water quality for consumption and agriculture purposes in recent period (Al Futaisi et al., 2007; Nag and Das, 2018; Kundu and Nag, 2018; Bhunia et al., 2018; Karunanidhi et al., 2020; Chakraborty et al., 2022; Chakraborty et al., 2021c; Sridhara et al., 2021; Gajalakshmi, 2022; Rathnamala and Shivashankara, 2022). Ground water quality data varies with time, area and methodology of sampling. Therefore, present values are differ from other workers carried in this area but their research work is more emphasize towards GIS. There is a research gap between our study and other researchers carried out in this area of Bagepalli taluk of Karnataka and hence, the present study is carried out. The water table has decreased in several parts of eastern Karnataka, despite the fact that rainfall has not decreased uniformly over the past three decades. While precipitation is diminishing in north inside Karnataka, it is expanding in south inside Karnataka. Water level depletion and yield dwindling in major districts, overexploitation of groundwater sources, and issues with water quality are major groundwater issues in the Bagepalli area. However, very few articles on water quality issues were published in the





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Bagepalli taluk. These factors justified to take the current study. The main objectives of the present study is to analyze the physico-chemical characteristics of water samples and assess the suitability of water for human consumption and irrigation. Also to find out the source of pollution. Create awareness about health hazards in drinking water. To understand the interrelationship of various chemical factors through statistical approaches. and recommendation for better management of ground water.

STUDY AREA

Bagepalli is a one of the Taluk in Chikballapur District (Figure 1) in the State of Karnataka, India. Bagepalli is situated 100 km north of Bangalore on the Bangalore - Hyderabad National Highway. The region is just below the southern border of the Rayalaseema in Andhra Pradesh, South India. It is semi-arid and drought prone with 560 mm of erratic and spatial rainfall. Bagapalli is located at 13.78°N 77.79°E. It has an average elevation of 707 metres (2319ft). The region is a semi arid drought prone one with low, erratic and spatial rainfall. The dust brown rocky terrain is severely undulating, with small hill ranges and outcrops that stud the topography. There is no mineral wealth and only a very thin and fragile soil cover. An adverse land: person ratio creates a strong thirst for cultivable land since less than one-half of the total land is fit for cultivation, with the remaining taken over by the hills and rocky fields. Hardly 5% of the cropped lands are irrigated by an age old network of rain-fed tanks (small lakes), each irrigating 2 to 10 hectares of wet land. The low water table is tapped through bore-wells drilled to more than 100 meters depth. Even these dry up in the summer months, from April to September every year, when temperatures rise to a dry heat of 38°C.

METHODOLOGY

A total of 10 water samples (S1 to S10) were collected from selected villages borewells of Bagepalli Taluk, Chikkaballapur District during June -2010 to May-2011. The temperature as recorded at the time of sampling by using mercury thermometer. The pH was measured with pH meter. All the other parameters were measured as per the Standard Methods (APHA,2005)

RESULTS AND DISCUSSION

Monthly fluctuation of physico-chemical parameters are given in the Table 4 to 11. Temperature of groundwater is largely dependent on atmospheric, heat, exothermic and endothermic reaction in rocks, infiltration of surface water, insulation, thermal conductivity of rocks, rate of movement of ground water and interference of man on the ground water. It is well know fact that the water temperature exerts direct and indirect influences on many abiotic and biotic components of any aquatic system (Das et al., 2000), During the study period, water temperature varied from a minimum of 25°C during Nov-2010 and a maximum of 35°C, May-2011 at all the sites (S 1 to S 10). In most of the natural water, the pH value is dependent on carbon dioxide-carbonate-bicarbonate equilibrium. As the equilibrium is markedly affected by temperature and pressure, it is obvious that changes in pH may occur when these are altered. Presence of phosphates silicates, borates, fluorides and other salts in dissociated form may also affect the pH. Most ground water has a pH range of 6 to 8.5. In the present study, pH concentration varied from a minimum of 7.0 during Jan-2011 at S7 to a maximum of 8.8 in Sept. 2010 at S1 sites, indicating circum neutral to alkaline nature of water. The recorded values are within stipulated limits for potable waters recommended as per CPCB norms (De Zwart and Trivedi,1995). pH has no direct adverse effects on health although values below 4.0 are known to produce some taste and higher values (>8.5) hasten the scale formation in water heating apparatus and also reduce the germicidal potential of chlorine. It is discernible from the above observation that the pH value are dependent on local soil conditions. The significant correlation of pH was observed with total hardness (r = -0.62, p < 0.06), magnesium hardness (r = 0.68, p < 0.06) and potassium (r = 0.6, p < 0.05). Electrical conductivity is the function of concentration of dissolved solids. Conductivity is the capacity of water to carry an electrical current and various both with number and types of ions, which in turn is related to the concentration





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of ionized substances in the water, most of the dissolved inorganic substances in water are in the ionized form and hence, contribute to conductance (Tiwari,2001). In present findings, the electrical conductivity values are found to be in the range of 302 to 2718 µmhos/cm (Table 5).EC showed direct relationship with total hardness, sulphate and total dissolved solids P<0.01). The total dissolved solids (TDS), value are found in the range of 311 to 2483 mg/L for ground water samples. That is well above the permissible limit (500ppm) of WHO. The maximum TDS value observed at site S 10 (2483 mg/L). The bulk of TDS include bicarbonates, sulphates and chlorides of calcium, magnesium, sodium and silica. High concentration of TDS increased water turbidity. Such water will have salty taste and produce scale on water heaters and cooking utensils (25). The total hardness (TH) values are found to be in the range of 311 (S 7) to 1042 (S 3) mg/L. Hardness of water was understood to be measure of the capacity of the water for precipitating soap. Soap is precipitated chiefly by the calcium and magnesium ions are usually present in significant concentration in natural waters. Hard water therefore is not suitable for domestic usage. Alkalinity in natural water is formed due to dissolution of carbon dioxide.

Carbonates and bicarbonates thus formed are dissociated to yield hydroxyl ions. The most common constituents of alkalinity in natural water include bicarbonates, carbonates and hydroxide. These compounds result from the dissolution of mineral substances in the soil, from waste water discharge, fertilizer and insecticides from agricultural runoff and microbial decomposition of organic material (24) . In the present study, the lowest value recorded 200 mg/L (S 5) and high value 588 mg/L (S 10). The highest value recorded in southwest monsoon and northeast monsoon and lower value recorded during summer season. Alkalinity showed positive correlation with calcium (p < 0.01, r = 0.61) and total hardness (p < 0.05, r = 0.65). Chlorides occur in all natural water in varying concentration. The chloride content normally increases as the mineral content increases. Most of the chloride in groundwater is present as sodium chloride. In the present study, chloride ranged from 80 mg/L (S 8) to 527 mg/L (S 10). Chloride showed a distinct seasonal variations and higher values were recorded during southwest monsoon and northeast monsoon season while lowest was recorded during summer season (Table-8). During the current study, the positive correlation was observed with total hardness (p < 0.05, r = 0.72) and magnesium (p < 0.05, r = 0.90). Nitrate enters the human body through the use of ground water for drinking and cause a number of health disorder, namely methemaglobinemia, gastric disorders, cancer, goiter, hypertension etc., when present in high concentration in drinking water. The main source of nitrate pollution is the ground water is use of more nitrogen fertilizer in the agricultural land. Many ground water have significant quantities of nitrate due to leaching of the nitrate with the percolation of water high amount of nitrate are generally indicative of pollution (26) . In the present investigation, the nitrate concentration varied between 48 mg/L (57) to 270 mg/L (S 2).

All the water samples are above permissible limit. The permissible limit is 45 mg/L as per the Indian standards, this is due to high use of fertilizer in agriculture field. Nitrate has showed a seasonal trend and noticed higher concentration during northeast and southeast monsoon season, while, in summer nitrate was recorded lesser then other season. In this study, nitrate showed significant positive correlation with sodium (p < 0.05, r = 0.79), potassium (p<0.05, r=0.76) and phosphate (p < 0.1, r = 0.92). Sulphate content in found water is made possible through oxidation, precipitation and constriction, as the water traverses through rocks. The source of sulphate in rocks are sulphur minerals, sulphides of heavy metals. Sulphide minerals, when oxidized give rise to soluble sulphates. In the present study, the sulphate concentration varied between 10 mg/L (S 5) to 95 (S 2) mg/L. Interestingly, the sulphate concentration were within the limits of Indian Standards (27). In the present study, sulphate showed a significant positive correlation with total iron (p < 0.05, r = 0.92), total hardness (p < 0.05, r = 0.66). Phosphorus in the natural fresh water is present mostly in inorganic forms, phosphorus being an important constituent of Biochemical systems may also be present in the organic form. The major source of phosphorus are domestic sewage, detergents, agricultural effluents with fertilizers and industrial waste waters. In the present study, the high phosphate value recorded 1.4 mg/L in S 5 and low value recorded 0.2 mg/L in S 1. During summer lower values recorded. Significant positive correlation was noticed with total hardness (p < 0.05, r =0.92), calcium hardness (p < 0.01, r = 0.89), magnesium hardness (p < 0.05, r = 0.92) and iron (p < -0.01, r = 0.65) (28)During the present investigation, maximum sodium was recorded to be 360 mg/L at S5 in the month of May 2011 and lowest value recorded 70 mg/L in the month of Dec 2010 at S 6. In summer it was recorded





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high when compared to other seasons. Most sodium salts are readily soluble in water, but take no part in chemical reactions, as do the salts of alkaline earth metals. For this reason sodium salt tend to remain in solution unless extracted during evaporation. In general these is a concomitant increases in sodium and chloride, the concentration of both increasing with total dissolved solids content (24). In the present study, sodium values showed a positive significant correlation was noticed with potassium p < 0.05, r = 0.70), fluoride < 0.05, r = 0.69), nitrate (p < 0.01, r = 78) and TDS (p < 0.05, r = 0.68). The common source of potassium are the silicate mineral, orthoclase, microline, nepheline, biotite igneous rocks and metamorphic rocks, etc., Although potassium is nearly as abundant as sodium in igneous rocks and metamorphic rock, its concentration in ground water is one tenth or even one-hundredth that of sodium. Two factors are responsible for the scarcity of potassium in groundwater one being the resistance to potassium minerals to decomposition by weathering The other fixation of potassium in clay minerals formed due to weathering. Potassium salts, being more soluble than sodium salts, are the last to crystallize during evaporation. In the present study, maximum potassium was recorded 4.2 mg/L at S 8 and minimum value recorded 1.2mg/L at S 3. In the present study a significant positive correlation was noticed with nitrate (p < 0.1, r = 0.82). Iron is one of the major constituents of rocks, next in abundance only to oxygen, silicon, and aluminium. The important iron-bearing minerals include pyroxens, amphiboles and micas among silicates, pyrite and chalcopyrite among sulphides and magnetite and hematite among oxides.

In igneous and metamorphic rocks, iron present mostly in the form of silicate minerals. Usually iron occurring in groundwater in the form of ferric hydroxide in concentration less than 0.5ppm. In the present investigation, the iron content varied between 0.1 mg/L (S 5) to 0.7 mg/L (S 3) (Table-11. All the iron values were within the permissible limits and no seasonal variation was observed in the present study. In the present investigation significant correlation was noticed with sulphate (p < 0.01, r = 0.68) and the Total dissolved solids (p < 0.05, r = 0.78). According to BIS the permissible limit of pH value for drinking water is 6.5 to 8.5. Abnormal values of pH in water causes bitter taste, affects mucous membrane, causes corrosion in pipelines and also affects aquatic life. The standard desirable limit of alkalinity in potable water is 200 mg/l as per BIS standards. Excess alkalinity in water is also harmful for irrigation which leads to soil damage by altering the soil pH which enhance soil pH to a great exert and reduce crop yields. Exceeding the permissible limits of hardness causes poor lathering with soap, deterioration of the quality of clothes, scale formation and skin irritation (Shashank Sourabh et al.,2014; Priyanka Khanna and Nidhi Rai,2016). According to Indian standards for drinking water, desirable limit of chloride is 250 mg/l, and the permissible limit is 1000 mg/l. Sulphate occurs due to discharge from gypsum and other minerals. Sulphate content in drinking water exceeding the 400 mg/L impart bitter taste and may cause gastro-intestine irritation and cantharsis (Priyanka Khanna and Nidhi Rai,2016; Manivasakam,2005) .

Thirumala and Kiran (2017) have analysed the ground water quality in Davangere town of Karnataka. Seasonal variations in the physico-chemical parameters in the ground water sample were observed by them and compared with BIS standards. They reported that ground water samples were moderately polluted and impact to health hazards. In their study, the water samples of all the 5 sites were quite good (fair) for irrigation purpose due to high salinity of ground water. The Geo spatial variations of ground water quality in Chickballapur District of Karnataka, India have been studied by Gajalakshmi and Anatharama (19) by utilizing the new technology. For their study, water samples were analyzed for 13 physico-chemical parameters and the results were compared with Indian standards. The geospatial distributions of ground water quality map of their study area have been prepared using spatial analyst tools in GIS. The spatial database established in GIS and results obtained in their study will be helpful for monitoring and managing ground water pollution. Sridhara et al (18) have studied the drinking water quality parameters in borewell samples by using GIS techniques in Chikkaballapura and Bagepalli taluks. They were suggested that in contaminated areas, continuous monitoring of ground water quality be carried out to stop further degradation and related concerns.





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One-way ANOVA with post-Hoc Tukey HSD Test

The p-value to the F-statistic of one-way ANOVA is lower than 0.05, suggesting that the one or more treatments are significantly different. These post-hoc tests would likely identify which of the pairs of treatments are significantly different from each other (Table1-3).

CONCLUSION

The present investigation has been made to evaluate the quality of ground water of Bagepalli Taluk area. A total of 10 underground water samples were collected from different locations. Water samples were analyzed for 15 physico-chemical parameters to evaluate their suitability for human consumption and domestic applications. In many samples the values of total hardness and chloride are exceed the permissible limits of WHO and BIS standards. It is concluded that the ground water quality is mainly affected by seepage of sewage, waste water and urban runoff into the ground waters, due to lack of proper sewerage facilities. Open drains are prevalent in study area. Further, lime nature of lithology and lack of rainfall due to the deforestation are also responsible for the degradation of ground water quality. Hence, it is recommended to the Karnataka state and Indian government to take care of ground water quality and to safeguard for the residents and tourists of Bagepalli Taluk, India. Thus to maintain the quality of ground water, preventive measures have to be taken to control the above parameters within the permissible limit. However, we could suggest that this water could be used for drinking after proper treatment (reverse osmosis etc.) and induced ground water recharge by construction of percolation tanks flooding of ground water by mixing surface water by promoting rainwater harvesting.

RECOMMENDATIONS AND SUGGESTIONS

Most of the bore-wells of in the study area contain high degree of carbonate hardness (Temporary). Hence, people must be advised to use boiled water for drinking. People must be educated regarding ill effects likely to be caused by the excess iron and bacteriological organisms. There should be a check on the use of phosphatic fertilizers as they also contribute for high fluoride concentrations which are being leached down to main groundwater body through irrigation practices. The residents in the village sector should be educated. An artificial recharge of groundwater may be adopted to reduce higher concentration of chemical parameters Further, concerned authorities should be directed to maintain the existing ponds, lakes, reservoirs in Megadi taluk to increase the ground water table. Some of the bore wells are liable to bacterial (E-Coil) contamination. Hence, for drinking purposes, water must be treated by boiling permanganate solution. Proper treatment of water is required before using the water for drinking purpose specially to reduce the iron content of water by the following methods. By adding alum, so that there will be sedimentation. By adding calculated amount of lime which is stirred properly so that the iron will be precipitated and separated. Restrict further drilling through bringing legislation and groundwater is treat as property of the State or Nation. Drip and sprinkler irrigation techniques must be adapted for water saving. Concerned district authorities should take necessary steps to create health awareness among the people with regard to the ill effects of fluorosis and water born diseases. Proper sanitation and drainage system should be provided to avoid contamination of drinking water. Practicing of environmentally compatible cropping patterns, bio-fertilizer and bio-pesticides should be encouraged. The present investigation has revealed that there is much scope for further study in the field of assessment of ground water recharge potential and estimation of pesticidal residues.

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Table 1: One way ANOVA data and independent treatment values

Treatment	A (pH)	B (NO3)	C(Total hardness)	D(Potassium)	Total
observations N	120	120	120	120	480





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sum	939.9000	13,998.1400	72,017.0000	314.1000	87,269.1400
mean	7.8325	116.6512	600.1417	2.6175	181.8107
sum of squares	7,378.2700	1,884,624.0196	47,433,347.0000	917.7900	49,326,267.0796
sample variance	0.1387	2,115.3333	35,402.8957	0.8036	69,853.4508
sample std. dev.	0.3724	45.9928	188.1566	0.8965	264.2980
std. dev. of mean	0.0340	4.1985	17.1763	0.0818	12.0635
Source	Sum of square	Degrees of freedom	Mean square	F statistic	P-value
treatment	28,995,021.5350	3	9,665,007.1783	1,030.4073	1.1102e-16
error	4,464,781.3856	476	9,379.7928		
total	33,459,802.9206	479			

Table 2: Tukey HSD and Scheffé data

Treatments pair	Tukey HSD Q statistic	Tukey HSD p- value	Tukey HSD inference		
A vs B	12.3083	0.0010053	** p<0.01		
A vs C	66.9950	0.0010053	** p<0.01		
A vs D	0.5899	0.8999947	insignificant		
Treatments pair	Tukey HSD Q statistic	Tukey HSD p- value	Tukey HSD inference		
B vs C	54.6867	0.0010053	** p<0.01		
B vs D	12.8981	0.0010053	** p<0.01		
C vs D	67.5849	0.0010053	** p<0.01		
	Scheffé TT-statistic	Scheffé p-value	Scheffé inference		





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A vs B	8.7033	3.5527e-15	** p<0.01
A vs C	47.3726	1.1102e-16	** p<0.01
A vs D	0.4171	0.9816553	insignificant
B vs C	38.6694	1.1102e-16	** p<0.01
B vs D	9.1204	1.1102e-16	** p<0.01
C vs D	47.7897	1.1102e-16	** p<0.01

Table 3: Bonferroni and Holm results: all pairs simultaneously compared

Treatments pair	Bonferroni & Holm T-statistic	Bonferroni p- value	Bonferroni inference	Holm p- value	Holm inference
A vs B	8.7033	0.0000e+00	** p<0.01	0.0000e+00	** p<0.01
A vs C	47.3726	0.0000e+00	** p<0.01	0.0000e+00	** p<0.01
A vs D	0.4171	4.0607888	insignificant	0.6767981	insignificant
B vs C	38.6694	0.0000e+00	** p<0.01	0.0000e+00	** p<0.01
B vs D	9.1204	0.0000e+00	** p<0.01	0.0000e+00	** p<0.01
C vs D	47.7897	0.0000e+00	** p<0.01	0.0000e+00	** p<0.01

Table 4. Monthly analysis of Water Temperature and pH in sites of some selected villages of Bagepalli Taluk

Sites	Jun e 201 0	Jul y 201 0	Augus t 2010	Sept. 2010	Oct. 2010	Nov 2010	Dec 201 0	Jan. 201 1	Feb 201 1	Mar 201 1	Apri 1 2011	Ma y 201 1
S1	33	31	30	27	28	25	26	27	30	31	32	35
S2	33	31	30	27	28	25	26	27	30	31	32	35
S3	33	31	30	27	28	25	26	27	30	31	32	35
S4	33	31	30	27	28	25	26	27	30	31	32	35
S5	33	31	30	27	28	25	26	27	30	31	32	35
S6	33	31	30	27	28	25	26	27	30	31	32	35
S7	33	31	30	27	28	25	26	27	30	31	32	35
S8	33	31	30	27	28	25	26	27	30	31	32	35



S

7 S

8 S

S10

8.2

8.1

8.3

8.2

8.1

8.0

8.1

8.1

7.9

7.8

8.0

7.9

7.7

7.6

7.9

7.

8



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7.0

7.1

7.1

7.1

7.3

7.3

7.4

7.3

7.3

7.6

7.6

7.5

8.1

7.9

7.9

7.9

8.2

8.1

8.1

8.2

7.7

7.2

7.3

7.3

59	33	31	30	27	28	25	26	27	30	31	32	33	
S10	33	31	30	27	28	25	26	27	30	31	32	35	
pH													
Sites	Jun e 201 0	Jul y 201 0	August 2010	Sept 2010	Oct 201 0	Nov. 2010	Dec. 2010	Jan. 2011	Feb 201 1	Mar. 2011	April 2011	May 2011	
S 1	7.8	7.1	8.1	8.8	7.4	7.6	7.6	7.4	7.4	7.8	7.6	7.6	
S 2	8.2	8.5	8.6	8.6	8.1	7.9	7.8	8.1	7.9	8.1	7.7	7.9	
S 3	7.7	8.0	8.1	8.6	7.7	7.8	7.9	7.4	7.5	8.2	7.9	7.7	
S 4	8.1	8.2	8.4	8.6	7.5	7.8	7.8	8.1	8.1	8.2	8.1	8.1	
S 5	7.9	8.2	8.1	8.3	7.8	7.6	7.6	7.5	7.9	7.5	7.6	7.6	
S 6	7.9	8.1	8.3	8.6	7.9	7.7	7.7	7.8	8.3	8.1	7.8	7.8	
			+	.						.			

Table 5. Monthly Electrical Conductivity (umhos/cm) and TDS (mg/L) in sampling sites of Bagepalli Taluk

7.6

7.5

7.7

7.6

7.4

7.3

7.5

7.4

Sites	June 2010	Jul y 201 0	Augus t 2010	Sept 2010	Oct. 201 0	Nov 2010	Dec. 2010	Jan. 201 1	Feb. 2011	Mar. 2011	Apri 1 2011	May 2011
S1	1900	2050	2100	2110	2122	2125	2100	1991	1786	1790	1802	1812
S2	720	1010	1020	1120	1140	1132	1202	1226	1043	1080	1025	1052
S3	1415	1495	1550	1535	1600	1632	1620	1642	1540	1311	1300	1381
S4	480	435	460	459	482	500	485	45 2	302	383	242	382
S5	1490	1545	1520	1531	1526	1640	1680	1542	1342	1311	1490	1346
S6	1490	1500	1520	155	1572	1620	1650	1650	1300	1352	1401	1451
S7	1137	1134	1132	1330	1128	1125	1123	1121	1106	1117	1128	1139
S8	1315	1312	1310	1307	1304	1302	1299	1296	1280	1292	1305	1318
S9	1125	1123	1120	11i8	1115	1112	1108	1105	1095	1105	1116	1127
S10	2718	2715	2712	2709	2704	2699	2693	2688	2642	2540	2550	2565



S8

S9

S10

S4



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Ramesha Iyyanahalli and Kiran TDS Jun Oct Dec Feb Mar Ma Jan. Augus Site July Sept. Nov. April y S1 S2 S3 S4 S5 S6 I 322 S7

Table 6 . Monthly analysis of Total hardness and Calcium hardness (mg/L) in different sites of some selected villages of Bagepalli Taluk

June 2010	July 2010	August 2010	Sept. 2010	Oct. 2010				Jan. 2011	Feb. 2011	Mar. 2011	April 2011	May 2011
711	752	864	762	657	594	1	620	636	511	543	512	597
582	572	672	745	642	583	3	622	663	522	498	422	387
665	792	842	972	968	913	1	1042	986	910	932	801	795
672	692	767	765	843	825	5	823	896	651	632	645	712
663	670	671	781	892	813	1	892	951	663	564	523	592
593	629	722	802	900	775	5	766	811	700	722	645	595
361	357	353	349	356	342	2	339	335	325	311	346	364
382	374	366	359	348	337	7	326	316	326	347	368	390
468	464	459	454	449	445	5	440	435	412	434	453	473
588	582	576	571	559	548	3	536	525	520	545	569	594
•				Calcium h	ardness	•						
June 2010	July 2010	August 2010	Sept. 2010	Oct. 2010	Nov. 2010			Jan. 2011	Feb. 2011	Mar. 2011	April 2011	May 2011
82	92	80	123	152	132	1	10	116	73	123	102	96
276	235	262	142	160	151	1	53	158	170	192	221	226
122	120	110	126	156	142	9	92	108	77	136	80	110
	2010 711 582 665 672 663 593 361 382 468 588 June 2010 82 276	2010 2010 711 752 582 572 665 792 672 692 663 670 593 629 361 357 382 374 468 464 588 582 June 2010 2010 82 92 276 235	2010 2010 2010 711 752 864 582 572 672 665 792 842 672 692 767 663 670 671 593 629 722 361 357 353 382 374 366 468 464 459 588 582 576 June 2010 August 2010 82 92 80 276 235 262	2010 2010 2010 2010 711 752 864 762 582 572 672 745 665 792 842 972 672 692 767 765 663 670 671 781 593 629 722 802 361 357 353 349 382 374 366 359 468 464 459 454 588 582 576 571 June 2010 July 2010 August 2010 Sept. 2010 82 92 80 123 276 235 262 142	2010 2010 2010 2010 2010 711 752 864 762 657 582 572 672 745 642 665 792 842 972 968 672 692 767 765 843 663 670 671 781 892 593 629 722 802 900 361 357 353 349 356 382 374 366 359 348 468 464 459 454 449 588 582 576 571 559 Calcium h June July August Sept. Oct. 2010 2010 2010 2010 2010 82 92 80 123 152 276 235 262 142 160	2010 2010 <th< td=""><td>2010 2010 2010 2010 2010 2010 2010 711 752 864 762 657 594 582 572 672 745 642 583 665 792 842 972 968 911 672 692 767 765 843 825 663 670 671 781 892 811 593 629 722 802 900 775 361 357 353 349 356 342 382 374 366 359 348 337 468 464 459 454 449 445 588 582 576 571 559 548 Calcium hardness June July 2010 2010 2010 2010 2010 2010 2010 2010 2010 2010 2010 2010 2010 2010<</td><td>2010 <th< td=""><td>2010 2010 2010 2010 2010 2010 2010 2010 2011 711 752 864 762 657 594 620 636 582 572 672 745 642 583 622 663 665 792 842 972 968 911 1042 986 672 692 767 765 843 825 823 896 663 670 671 781 892 811 892 951 593 629 722 802 900 775 766 811 361 357 353 349 356 342 339 335 382 374 366 359 348 337 326 316 468 464 459 454 449 445 440 435 588 582 576 571 559 548</td><td>2010 2010 2010 2010 2010 2010 2010 2011 2011 2011 711 752 864 762 657 594 620 636 511 582 572 672 745 642 583 622 663 522 665 792 842 972 968 911 1042 986 910 672 692 767 765 843 825 823 896 651 663 670 671 781 892 811 892 951 663 593 629 722 802 900 775 766 811 700 361 357 353 349 356 342 339 335 325 382 374 366 359 348 337 326 316 326 468 464 459 454 449 445</td><td>2010 2010 2010 2010 2010 2010 2011 <th< td=""><td>2010 2010 2010 2010 2010 2010 2011 <th< td=""></th<></td></th<></td></th<></td></th<>	2010 2010 2010 2010 2010 2010 2010 711 752 864 762 657 594 582 572 672 745 642 583 665 792 842 972 968 911 672 692 767 765 843 825 663 670 671 781 892 811 593 629 722 802 900 775 361 357 353 349 356 342 382 374 366 359 348 337 468 464 459 454 449 445 588 582 576 571 559 548 Calcium hardness June July 2010 2010 2010 2010 2010 2010 2010 2010 2010 2010 2010 2010 2010 2010<	2010 2010 <th< td=""><td>2010 2010 2010 2010 2010 2010 2010 2010 2011 711 752 864 762 657 594 620 636 582 572 672 745 642 583 622 663 665 792 842 972 968 911 1042 986 672 692 767 765 843 825 823 896 663 670 671 781 892 811 892 951 593 629 722 802 900 775 766 811 361 357 353 349 356 342 339 335 382 374 366 359 348 337 326 316 468 464 459 454 449 445 440 435 588 582 576 571 559 548</td><td>2010 2010 2010 2010 2010 2010 2010 2011 2011 2011 711 752 864 762 657 594 620 636 511 582 572 672 745 642 583 622 663 522 665 792 842 972 968 911 1042 986 910 672 692 767 765 843 825 823 896 651 663 670 671 781 892 811 892 951 663 593 629 722 802 900 775 766 811 700 361 357 353 349 356 342 339 335 325 382 374 366 359 348 337 326 316 326 468 464 459 454 449 445</td><td>2010 2010 2010 2010 2010 2010 2011 <th< td=""><td>2010 2010 2010 2010 2010 2010 2011 <th< td=""></th<></td></th<></td></th<>	2010 2010 2010 2010 2010 2010 2010 2010 2011 711 752 864 762 657 594 620 636 582 572 672 745 642 583 622 663 665 792 842 972 968 911 1042 986 672 692 767 765 843 825 823 896 663 670 671 781 892 811 892 951 593 629 722 802 900 775 766 811 361 357 353 349 356 342 339 335 382 374 366 359 348 337 326 316 468 464 459 454 449 445 440 435 588 582 576 571 559 548	2010 2010 2010 2010 2010 2010 2010 2011 2011 2011 711 752 864 762 657 594 620 636 511 582 572 672 745 642 583 622 663 522 665 792 842 972 968 911 1042 986 910 672 692 767 765 843 825 823 896 651 663 670 671 781 892 811 892 951 663 593 629 722 802 900 775 766 811 700 361 357 353 349 356 342 339 335 325 382 374 366 359 348 337 326 316 326 468 464 459 454 449 445	2010 2010 2010 2010 2010 2010 2011 <th< td=""><td>2010 2010 2010 2010 2010 2010 2011 <th< td=""></th<></td></th<>	2010 2010 2010 2010 2010 2010 2011 <th< td=""></th<>





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S5	231	241	212	217	196	245	289	195	242	132	232	156
S6	75	90	80	251	260	282	310	312	157	119	78	132
S7	54	53	52	52	51	51	50	50	49	47	49	52
S8	55	54	53	52	50	50	49	48	48	49	51	56
S9	35	33	32	31	32	32	31	30	32	34	37	38
S10	49	48	47	46	46	45	45	43	44	45	47	50

Table 7: .Monthly analysis of Magnesium Hardness and Total alkalinity(mg/L) in sampling sites of selected villages of Bagepalli Taluk

Sites	June 2010	July 2010	August 2010	Sept. 2010	Oct. 2010	Nov. 2010	Dec 2010		Jan. 2011		eb. 111	Mar. 2011	April 2011	May 2011
S1	31	32	33	32	41	50	51		62	3	0	32	29	30
S2	39	37	38	39	39	59	40		42	4	0	33	25	30
S3	31	35	37	32	38	40	41		46	2	2	20	18	23
S4	31	36	39	38	41	36	44		43	3	3	32	31	32
S5	39	37	39	38	40	38	42		52	5	7	34	32	32
S6	33	41	44	44	49	59	41		42	4	4	35	32	32
S7	52	53	53	52	51	50	49		49	4	7	49	51	53
S8	58	57	51	54	53	53	52		52	5	1	53	54	58
S9	38	37	36	36	35	34	34		33	3	3	34	36	38
S10	90	89	89	88	86	84	82		81	8	0	84	87	89
					To	otal alka	linity							
Sites	June	July	August	Sept.	Oc		Nov.	De		Jan.	Feb.	Mar.	April	May
	2010	2010	2010	2010	201	10	2010	201	10	2011	2011	2011	2011	2011
S1	375	370	410	380	3	85	390	36	60	300	350	375	280	320
S2	300	365	380	390	4	.00	335	31	10	320	340	360	290	310
S3	300	310	360	400	3	80	375	37	75	280	330	380	310	360
S4	300	340	350	410	4	20	350	30	00	400	240	280	225	260
S5	280	310	410	460	4	.00	420	44	40	375	200	250	260	265
S6	375	400	415	420	4	80	490	49	95	390	375	395	360	320
S7	433	429	424	420	4	11	403	39	95	386	374	395	416	437
S8	483	478	473	468	4	.63	459	45	54	449	427	447	467	488
S9	484	476	474	469	4	.60	450	44	41	432	418	442	465	489
S10	588	582	576	570	5	58	547	53	36	524	520	544	569	572

Table 8 Monthly Chloride and Nitrate (mg/L) in various sampling sites of Bagepalli Taluk

Sites	June	July	August	Sept.	Oct.	Nov.	Dec.	Jan.	Feb.	Mar.	April	May
	2010	2010	2010	2010	2010	2010	2010	2011	2011	2011	2011	2011



S10



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Ramesha Iyyanahalli and Kiran S1 S2 S3 .460 S4 S5 S6 S7 S8 S9 S10 Nitrate Sites June July August Sept. Oct. Nov. Dec. Jan. Feb. Mar. April May S1 S2 S3 S4 .140 S5 S6 S7 S8 S9

Table 9. Monthly data of Sulphate and phosphate (mg/L) in sites of some selected villages of Bagepalli Taluk

Sites	June	July	August	Sept.	Oct.	Nov.	Dec.	Jan.	Feb.	Mar.	April	May
	2010	2010	2010	2010	2010	2010	2010	2011	2011	2011	2011	2011
S1	47	43	42	50	47	43	52	52	57	53	52	53
S2	78	70	74	70	79	81	82	85	90	94	92	95
S3	43	42	50	47	43	52	57	55	53	52	47	48
S4	48	46	46	47	47	34	36	34	35	49	44	46
S5	15	12	11	10	10	16	15	10	13	10	11	11
S6	27	24	22	28	25	22	29	40	40	39	39	40
S7	46	46	47	47	34	36	34	35	49	44	46	40
S8	65	67	66	66	52	57	50	45	58	57	53	57
S9	25	23	24	20	30	31	34	40	25	21	20	24
S10	48	43	41	48	45	42	50	56	53	51	48	48
	•	•	•	•	Phospha	ate		•			•	•





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Sites	June	July	August	Sept.	Oct.	Nov.	Dec.	Jan.	Feb.	Mar.	April	May
	2010	2010	2010	2010	2010	2010	2010	2011	2011	2011	2011	2011
S1	0.3	0.3	0.4	0.4	0.5	0.4	0.5	0.6	0.2	0.2	0.3	0.2
S2	0.7	0.8	0.9	0.9	1.0	1.1	1.2	1.3	1.2	1.2	1.0	0.9
S3	0.6	0.9	0.9	1.0	0.9	0.4	0.2	0.3	0.2	0.3	0.4	0.8
S4	0.9	1.0	0.9	1.1	0.9	0.9	1.0	1.3	0.4	0.5	0.4	0.7
S5	1.2	1.1	1.3	1.0	1.4	1.4	1.2	1.1	0.9	0.8	0.8	1.1
S6	1.1	1.0	1.0	0.9	0.9	1.0	1.1	1.2	0.2	0.3	0.4	0.8
S7	0.9	1.0	0.9	1.1	0.9	1.0	1.3	0.4	0.5	0.5	0.4	0.9
S8	0.8	0.8	0.9	0.7	0.9	0.9	1.2	1.3	0.9	0.8	1.0	0.9
S9	1.0	0.9	0.8	1.0	0.8	0.8	1.0	1.1	0.3	0.5	0.4	0.5
S10	1.2	1.3	1.2	1.4	1.2	1.2	1.3	1.2	0.9	0.8	0.7	1.0

Table 10 . Monthly analysis of Sodium and Potassium (mg/L) in sites of some selected villages of Bagepalli Taluk

Sites	June 2010	July 2010	August 2010	Sept. 2010	Oct. 2010	Nov. 2010	Dec. 2010	Jan. 2011	Feb. 2011	Mar. 2011	April 2011	May 2011
S1	190	185	180	185	210	215	220	240	260	245	240	210
S2	150	145	140	145	175	180	185	210	225	210	215	195
S3	200	210	190	180	160	165	150	175	235	215	240	260
S4	240	235	215	215	245	235	205	210	245	240	260	280
S5	245	265	260	265	280	260	280	300	340	340	345	360
S6	81	80	77	76	74	72	70	69	72	75	78	82
S7	112	110	107	105	103	100	98	96	100	105	109	114
S8	133	131	128	125	122	120	117	114	119	125	130	136
S9	186	183	179	175	171	167	136	160	166	174	182	190
S10	145	142	139	136	134	131	128	125	130	136	142	148
1												

Potassium

					100000							
Sites	June 2010	July 2010	Augus t 2010	Sept	Oct. 2010	Nov	Dec. 2010	Jan. 2011	Feb. 2011	Mar. 2011	Apri	May 2011
Sites	2010	2010	1 2010	2010	2010	2010	2010	2011	2011	2011	1 2011	2011
S1	4.1	4.1	4.0	3.9	3.9	3.8	3.7	3.7	3.8	3.9	4.0	4.1
S2	3.0	3.0	3.1	3.1	3.0	3.0	2.9	2.9	3.0	3.1	3.2	3.3
S3	1.4	1.3	1.3	1.2	1.2	1.3	1.2	1.2	1.3	1.4	1.4	1.5
S4	3.0	3.0	4.0	4.0	3.0	3.0	3.1	3.1	3.0	3.1	4.0	4.0
S5	2.0	2.1	2.1	2.1	3.0	3.0	3.1	2.8	2.8	3.0	3.2	3.4
S6	1.6	1.5	1.5	1.4	1.4	1.3	1.3	1.2	1.3	1.5	1.6	1.7
S7	2.1	2.1	2.0	2.0	1.9	2.0	2.0	1.8	1.9	2.0	2.1	2.2
S8	3.0	3.0	3.0	4.0	4.0	3.0	3.0	3.0	3.0	4.0	4.1	4.2
S9	2.0	2.0	2.0	1.9	1.9	1.8	1.8	1.9	1.9	2.0	2.1	2.1





S10	3.1	3.1	3.0	3.0	2.9	2.9	2.8	2.8	2.9	3.0	3.1	3.2

Table 11: Monthly Iron (mg/L) in sampling sites of Bagepalli Taluk

Sites	June 2010	July 2010	August 2010	Sept. 2010	Oct. 2010	Nov. 2010	Dec. 2010	Jan. 2011	Feb. 2011	Mar. 2011	April 2011	May 2011
S1	0.5	0.5	0.5	0.6	0.5	0.5	0.5	0.4	0.4	0.4	0.3	0.3
S2	0.6	0.4	0.4	0.3	0.3	0.4	0.4	0.2	0.3	0.3	0.4	0.5
S3	0.7	0.6	0.5	0.5	0.5	0.6	0.6	0.6	0.5	0.6	0.5	0.4
S4	0.3	0.3	0.3	0.4	0.4	0.2	0.3	0.2	0.2	0.2	0.3	0.2
S5	0.4	0.4	0.3	0.3	0.3	0.2	0.3	0.3	0.2	0.3	0.4	0.1
S6	0.4	0.4	0.4	0.3	0.5	0.3	0.4	0.3	0.3	0.3	0.2	0.3
S7	0.4	0.3	0.3	0.2	0.4	0.3	0.4	0.3	0.3	0.2	0.2	0.3
S8	0.5	0.5	0.4	0.6	0.5	0.5	0.5	0.4	0.4	0.4	0.3	0.3
S9	0.4	0.4	0.3	0.3	0.3	0.2	0.4	0.4	0.3	0.2	0.2	0.3
S10	0.5	0.5	0.4	0.4	0.5	0.5	0.5	0.4	0.4	0.3	0.3	0.2

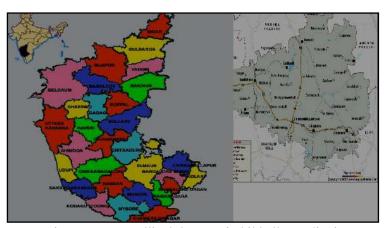


Figure 1: Bageapalli taluk map of Chikballapur district





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RESEARCH ARTICLE

Impacts of Microplastic Pollution in the Nanjarayan Lake - an Issue of Concern

M. Nandhini^{1*} and. J.Jagatheeswari²

¹Research Scholar, Department of Zoology, Chikkanna Government Arts College, Tiruppur, (Affiliated to Bharathiar University, Coimbatore), Tamil Nadu, India.

²Assistant Professor, Department of Zoology, Chikkanna Government Arts College, Tiruppur, (Affiliated to Bharathiar University, Coimbatore), Tamil Nadu, India.

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*Address for Correspondence

M. Nandhini

Research Scholar, Department of Zoology, Chikkanna Government Arts College, Tiruppur, (Affiliated to Bharathiar University, Coimbatore), Tamil Nadu, India.

Email: mbbsnandhu@gmail.com



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ABSTRACT

This study examines the widespread usage of plastics and the growing body of evidence on its possible eco-toxicological effects, particularly on smaller plastic particles known as micro-plastics. It addresses developing alternatives and their environmental suitability, as well as potential mitigation techniques aimed at reducing the prevalence of (micro)plastics. This paper evaluates these existing tools, examines their scientific foundations, and explores potential roadblocks to the applicability and relevance of current and upcoming legislative initiatives. The Nanjanarayan Lake, which has been preserved as a bird sanctuary, will also be the subject of a more focused investigation.

Keywords: Micro-plastics; Nanjanarayan lake; Rising concern; Toxic impacts; Environmental impacts.

INTRODUCTION

The name "plastic" serves as a catch-all for a variety of products manufactured of synthetic or semi-synthetic organic molecules. Plastics are described as "polymeric materials that may contain other substances to improve performance and/or reduce costs" by the International Union of Pure and Applied Chemistry (IUPAC). These very malleable materials are capable of being molded into solid things in a wide variety of sizes and shapes. (Baztan et.al., 2017) Plastics are polymers, which are made up of long chains of linked, identical units (sometimes known as "monomers"). Imagine a polymer as being similar to a pearl necklace in which the monomers are the individual





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pearls as one way to see this. Polymerization is the process by which these monomers are joined, and as a result, plastics can be categorized based on the chemical process used to create them, such as condensation, poly-addition, or cross-linking, or based on the chemical make-up of the polymer's backbone and side chains. The silicones, acrylics, polyesters, polyurethanes, and halogenated plastics subcategories are among the most significant. (Thompson, 2019) Primary micro-plastics, which are made on purpose to be millimeter or sub-millimeter sized, are present in a wide range of everyday items, including toothpaste, exfoliating lotions, facial cleansers, and other personal hygiene products. These products are particularly concerning because, according to estimates, 6% of all liquid skin-cleansing solutions marketed for sale in the EU, Switzerland, and Norway include micro-plastics, the majority of which are made of the plastic polyethylene (PE). (Walker *et.al.*, 2018)

The raw materials used to make plastic products are a significant source of primary micro-plastics. Primary micro-plastics may accumulate owing to improper handling, unintentional loss, runoff from processing facilities, and product residues. Micro-plastics are utilized in air-blasting media and, to a lesser extent, in medicine, namely as drug delivery systems. Micro-plastics are released into home waste waters after being used, where they may end up in the environment. In order to limit the amount of primary micro-plastics entering the environment, it is possible to identify their sources as well as their precise origins. (Zalasiewicz *et.al.*, 2015)

Environmental impacts of plastics and microplastics

Effects of plastics and micro-plastics on the environment "End of life" does not necessarily mean "end of impact". In reality, it has become obvious that there is no such thing as "end of life" for plastics because these materials continue to exist and pollute long after they have served their intended purpose. Depending on how it is used, plastic may, when it reaches the trash stage of its life cycle, constitute a serious threat to the environment and the climate. A 2019 analysis states that by 2015, the handling of plastic garbage worldwide looked like Figure. (Grinevald, 2011) According to the most recent statistics, 7.2 million tonnes of plastic post-consumer trash in Europe (both inside and outside the EU) was land filled, 9.4 million tonnes was collected for recycling, and roughly 12.4 tonnes was burned. (Vert *et.al.*, 2012)

The fate of plastics in the environment

It is intrinsically difficult to predict how (micro)(nano)plastics will behave in the environment. This is mostly caused by the variety of sources and entry points into the environment as well as the length of time required to ascertain the pathways for their breakdown. This also applies to smaller particles because of their size (see Figure). As a result, it can be challenging to quantify these materials, especially given the lack of established techniques for sampling, unit normalization, data expression and quantification, and identification, especially for smaller-sized plastics. Additionally, these materials lack a common definition, particularly in the case of nanoplastics. Micro-plastics have been found all over the world, especially in remote areas, from the Arctic to the Antarctic, and all the way down in the water column (benthos). However, micro-plastics are also discovered in agricultural soils, sediments, and even the atmosphere, both inside and outside of buildings. They are also discovered in rivers and lakes. It demonstrates the variety of routes through which plastics are introduced to the ecosystem, particularly the marine environment. ()

Plastics can degrade once they are in the environment through biotic and/or abiotic processes. The former comes before the latter and is a crucial initial step. In other words, abiotic degradation processes must come before biodegradation mechanisms. As a result, particles have greater surface area-to-volume ratios and are less stable structurally and mechanically, making them more accessible to microbial action. (Kricheldorf, 2011)

EFFECTS OF PLASTICS

One of the main perceived dangers to biodiversity is plastic pollution. It is particularly concerning due to its prevalence, toughness, and persistence in the environment. More than 90% of all contacts between garbage and people in the waters involve plastic debris. At least 17% of the species impacted by entanglement and ingestion were identified as threatened or near threatened by comparing the listed encounters with the International Union for Conservation of Nature (IUCN) Red List. The interaction of living things with plastic waste has a wide range of direct





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and indirect impacts, including the possibility of sub-lethal effects, which, due to their unpredictability, may be of great concern. In general, the presence of bigger plastic items in the ocean may cause entanglement and ingestion, the possible development of new habitats, and raft-based dispersal, including the transportation of invasive species. Although the obtained data seems to indicate that entanglement is significantly more lethal (79% of all instances) than ingestion (4% of all cases), both entanglement and ingestion commonly result in damage or death. (da Costa et.al., 2016)

Debris may also create new habitats, and abandoned fishing gear, for instance, has been found to kill invertebrates through "ghost fishing" as well as create new habitats for them. In recent decades, there has been a rise in the dispersal of species in the maritime environment, particularly species without a pelagic larval stage. Numerous species that are highly dependent on ocean currents have always built rafts out of natural materials like wood, but industrialization and the steady growth in the amount of plastic trash in the oceans suggest that rafting is actively contributing to the dispersal of this debris. (Barnes *et.al.*, 2019) It was interesting to see that the uptake of microplastics depends not only on their size and shape but also, and maybe less logically, on their color, with yellow particles being preferentially ingested. They resemble prey, which is probably why. Ingestion of micro-plastics can directly cause internal damage and digestive tract obstruction, which commonly result in lower food intake and thus reduced nutrition. Potentially, this brings about famine and death. Micro-plastics have been shown to lodge in the gills of air-breathing creatures, which may result in slower respiration rates. There are still few studies examining how these extremely widespread compounds affect terrestrial environments. The characteristics that make up the biota are the same regardless of how different soils and aquatic habitats are from one another because many organisms flourish in small bodies of water that are present at or just below the surface, making them basically aquatic species. (Jambeck *et.al.*, 2015)

Additionally, earthworms and mites have the ability to consume micro-plastics, which most likely contributes to their existence and accumulation in food webs. For instance, it was found that the earthworm *Lumbricus terrestris* had its growth rate significantly reduced, and that its mortality rates had increased. Additionally, these earthworms transferred micro-plastics from the litter into the soil by size-selectively transporting them downward through their tunnels. Additionally, it was found that only the smaller particles that the earthworms were exposed to were digested. Given how heavily earthworms influence the physical characteristics of soil, this finding could have significant ramifications for the fate and risk of micro-plastics in terrestrial ecosystems. (da Costa, 2014)

Regardless of the materials employed, these plastics nevertheless share some of the problems with conventional plastics, as many are only "biodegradable" under certain circumstances, such as prolonged high humidity and high temperatures. Such circumstances are infrequently possible in conventional composting facilities, making them unsuitable for home composting. This also means that these purportedly biodegradable materials take far longer to disintegrate after being dumped than expected. Claims about biodegradablity need to be carefully examined. Some substances that were once marketed as "biodegradable," "oxo-degradable," or "oxo-biodegradable" had an oxidizing ingredient that aided in the substance's breakdown in the presence of oxygen, UV radiation, and heat. The final outcome, nevertheless, was the production of minute plastic pieces. Furthermore, these goods are not suitable for long-term reusable applications because of the way they are made, which causes them to begin to fragment in a short amount of time. Due to the oxidizing agent's presence, such items could not be composted either, as doing so might cause the compost's quality and market value to decline. The European Commission has suggested that actions be taken against certain materials across the EU as a result of a report. (Zhang et.al., 2010)

Biodegrading organisms

From an environmental perspective perhaps the most urgent question is what to do with the vast amounts of plastic already found in the environment. For larger debris, multiple actions exist, including increasing citizen participation in clean-up actions. However, they have proven to be insufficient. Hence, research has focused on the development of biotechnology-based strategies centred on the process of bio-degradation. (Gouin *et.al.*, 2015)





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The development and use of plastic-consuming organisms will however not contribute to a reduction in the significant volume of greenhouse gas emissions that occur throughout the plastic life-cycle. Multiple bacteria and fungi have been reported as potential tools in the bio-degradation of plastics, including the bacterial species *Pseudomonas aeruginosa*, *P. stutzeri*, *Streptomyces badius*, *S. setonii*, *Rhodococcus ruber*, *Comamonas acidovorans*, *Clostridium thermocellum and Butyrivibrio fibrisolvens*. (Zettler *et.al.*, 2013)

Fungal strains described as viable biotechnological agents in the bioremediation of plastics include *Aspergillus niger, A. flavus, Fusarium lini, Pycnoporus cinnabarinusand and Mucor rouxii*. These organisms have been isolated from diverse environments, including landfills, plastic surfaces buried in soils, marine water or mangrove soil. Demonstrated within laboratorial settings, it is important to note that the reported efficiency of such biotechnological-based approaches is obtained under optimised conditions for organisms and substrates. This efficiency may be enhanced by, for example, subjecting plastics to different chemical and/or physical pre-treatments. Treatments include photolysis mediated by UV radiation and ozone and thermal treatments, but the effectiveness of the treatments greatly depends on the nature of the polymer. (Lebreton et.al., 2018)

However, because of the inherent expenses, such per-treatments are improbable procedures in potentially large-scale operations. Along with the price of the actual treatments, there are expenses associated with gathering and managing the plastic materials that need to be treated. As a result, the outcomes ought to be viewed as proof of concept. Additionally, it should be emphasized that using these biodegrading organisms in the environment directly is improbable or simply not practicable considering the accompanying dangers and uncertainties for ecosystems as well as for human health. (Ayre, 2016)

CONCLUSIONS

The presence, fluxes, paths, fates, and consequences of plastics in many environmental compartments have yet to be consistently and reliably measured by research. Marine pollution has received far more attention, both because of the immediate attention generated by the pollution's effects on biota that can be seen directly and because of the plastics' demonstrably widespread and transnational presence. However, other estimations opine that contamination levels in freshwater systems and soils may be higher than those reported for the marine environment, particularly in the case of micro-plastics. Nevertheless, despite these well-informed hypotheses, there is still a great deal of speculation and numerous known unknowns. These uncertainties and knowledge gaps substantially impair a full assessment of the health implications and limit an educated decision by consumers, communities and policy makers. At every stage of the life-cycle of plastic materials, inadequate and incomplete understanding also contributes to possibly long-term negative effects on the environment and human health. Therefore, it is vital to attract more concentrated scientific and governmental attention to these environmental compartments, not at the expense of, but in addition to the current marine (micro)plastic pollution studies.

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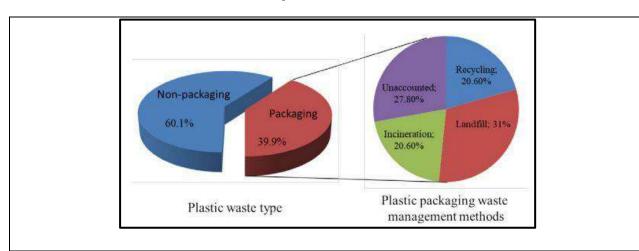


Figure 1. Global plastic waste management, Vert et.al., 2012.



Figure 2. Microplastics affecting animals (Jambeck et.al., 2015)





RESEARCH ARTICLE

The Effect of Whole-Body Vibration and Mulligan Mobilization with Movement in the Reduction of Pain and Disability in Subjects with **Knee Osteoarthritis**

P. Saranraj^{1*}, S. Rajadurai² and P. Susmitha³ B. Arun⁴, P. Savitha⁵

¹Ph.D Scholar, Department of Physiotherapy, Meenakshi Academy of Higher Education and Research Institute, Chennai, Tamil Nadu, India. Professor, Nandha college of Physiotherapy, (Affiliated to Tamil Nadu Dr. MGR Medical University). Erode, Tamil Nadu, India.

²M.S. Ortho, Department of Orthopaedics, Meenakshi Medical College Hospital and Research Institute, Kanchipuram, Tamil Nadu, India.

³Associate Professor, Department of Cardio Respiratory, Nandha College of Physiotherapy (Affiliated to Tamil Nadu Dr. MGR Medical University), Erode, Tamil Nadu, India.

⁴Grade II Physiotherapist, Thanthai Periyar Government Headquarters Hospital, Erode, Tamil Nadu, India.

⁵BPT internee Nandha college of Physiotherapy, Erode, Tamil Nadu, India.

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*Address for Correspondence

P. Saranraj

Ph.D Scholar,

Department of Physiotherapy,

Meenakshi Academy of Higher Education and Research Institute,

Chennai, Tamil Nadu, India.

Professor,

Nandha college of Physiotherapy,

(Affiliated to Tamil Nadu Dr. MGR Medical University).

Erode, Tamil Nadu, India. Email: saranraj.pt@gmail.com



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ABSTRACT

Knee osteoarthritis also known as the degenerative joint disease of the knee, is a common chronic progressive joint disease characterized by articular cartilage destruction and degeneration, adjacent subchondral bone lesions, bone hyperplasia and osteophyte formations at the joint margin. The aim of this study to compare the effect of whole-body vibration therapy and mulligan's mobilization with movement forknee osteoarthritis. A experimental study design consisting of 30 patients with knee osteoarthritis divided into two group at the average age was about 45 to 55 years. Group A underwent Whole Body vibration and Group B underwent Mulligan's Mobilization with movement. All the patients underwent pre and post-test scores of VAS and WOMAC. The calculated t-test values by the unpaired





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test in group A and group B are 7.4 and 12.8. In the present sample, both groups show positive effect but mulligan's mobilization with movement group has a superior effect than whole body vibration group.

Keywords: whole-body Vibration, mulligan's Mobilization with movement, VAS, WOMAC

INTRODUCTION

Knee osteoarthritis, sometimes referred to as the degenerative joint disease of the knee, is a common, chronic, and progressive joint condition marked by articular cartilage degradation and degeneration, nearby subchondral bone lesions, bone hyperplasia, and osteophyte forms at the joint boundary. Primary and secondary osteoarthritis are the two categories into which osteoarthritis may be divided. Ageing is a factor in primary osteoarthritis, a persistent degenerative condition.[1] As cartilage ages, its water content diminishes, making it more susceptible to breakdown. Contrarily, secondary osteoarthritis often develops in the early stages of life as a result of specific circumstances such as an accident, extended amounts of time spent stooping or crouching at work, diabetes, and obesity. The prevalence of knee osteoarthritis was determined to be 28.7% nationwide. It primarily affects older people. There are primary and secondary forms of osteoarthritis.[2] By 2020, osteoarthritis will rank as the fourth leading cause of disability, according to the World Health Organization. According to reports, OA has an economic, psychological, and physical toll. According to a World Health Organization study, 80 per cent of osteoarthritis patients over 60 have reduced mobility, and osteoarthritis is the main cause of disability worldwide[3]. Around 13% of women and 10% of men 60 years of age and older have symptoms of knee osteoarthritis. Osteoarthritis of the knee affects bodily functions including muscular strength, proprioception, and joint stability as well as causes joint pain and stiffness[4].

The aberrant mechanical stress brought on by joint instability is one of the main elements leading to the development and progression of knee osteoarthritis, according to reports on the pathogenesis of the condition. Agonists and antagonists cause disruptions that affect the dynamic and static stability of the knee joint, which results in the etiopathogenesis of knee pain. The tibial and femoral articular surfaces of the knee joint are anatomically unstable, and the surrounding ligaments and menisci keep the joint stable. Joint instability is often brought on by functional anomalies in these tissues[5]. A modern method of joint mobilisation known as Mulligan's notion of mobilisation with movement (MWM) involves a therapist using an auxiliary gliding force that is pain-free while still including active movement. The idea behind mobilisation with movement is to correct the postural defect and restore the knee's biomechanics. The quick recovery of pain-free movement may be attributed to the combination of joint mobilisation and active movement. A decrease in pain is said to be the outcome of correcting positional faults since the processes behind the effectiveness of mobilisation with movement are based on mechanical dysfunction[6].

This concept is related to positional faults that occur due to injury or changes in the shape of articular surfaces, the thickness of cartilage, orientation of fibres of ligaments and capsules which lead to mal tracking of the joints A recent systemic review heightened the problem that mobilisation with movement exhibits the immediate effects on pain and disability. This technique aims at restoring pain-free and a full range of motion in the joint[7] Whole Body Vibration Therapy (WBVT) is a feasible and curative strength exercise technique that has received considerable attention in recent years. A vibrating plate produces vibrations, which are transferred from surfaces that come into touch with the body to activate muscles and tendons. Whole-body vibration treatment has been shown in trials to enhance neuromuscular function and provide an aged population with a safe programme. It is a quick and secure procedure for treating persistent osteoarthritis of the knee[8]. Whole-body vibration exercise may be used to increase physical functions, reduce pain, improve neuromuscular function and enhance the multi joint strength performance of lower limbs during a counter-movement jump. Exercise and physical activity were suggested as the main treatments for knee osteoarthritis by the UK's National Institute, of Health and Clinic Excellence. It improves the lower extremities' multi joint strength and muscular strength. Whole body vibration may be helpful for people with





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knee osteoarthritis because it can reduce pain, enhance balance control, and improve gait pattern, and because basic balancing ability and mobility are substantially connected with the improvement of knee-specific function[9].

METHODS

PARTICIPATION

A total of 30 patients were included in this study all subjects provided written informed consent before entering the study. The patient's age is between 45 years to 55 years. Inclusion criteria were as follows: Patients with osteoarthritis knee pain.

STUDY DESIGN

Quasi-Experimental study design

OUTCOME MEASURES

VISUAL ANALOG SCALE (VAS) is the scoring test that is a measurement of pain that was given at any time of day. HE WESTERN ONTARIO AND MCMASTER UNIVERSITIES OSTEOARTHRITIS INDEX (WOMAC) is the scoring test that is a measurement of disability that was given at any time of day.

INTERVENTION

For 12 weeks, 15 patients in group A had whole-body vibration therapy three times per week with at least one day in between sessions. In this group B, 15 patients had mulligan mobilisation with movement. For a period of six days per week, the mobilisation with movement (MWM) approach was used with three sets of ten repetitions on each treatment occasion. Performing the workouts minimises muscular soreness, impairment, discomfort, and increases function. Under the direction of the therapist, each subject is shown how to do each exercise. The patient is instructed in the following interventions as part of a home programme. Additionally, patients are routinely observed to determine the correctness of the findings.

Training Program

Group A

Exercises included

The participants will stand with slightly bent knees (30°) on the platform and without shoes. To prevent the feeling of uncomforted and anxious, the patient can hold the handle during training

Group B

Exercises included
MULLIGAN MOBILIZATION FOR MEDIAL GLIDE:
MULLIGAN MOBILIZATION FOR LATERAL GLIDE
MULLIGAN MOBILISATION FOR ROTATION GLIDE

STATISTICAL ANALYSIS

The data were evaluated by using an unpaired 't' test. The unpaired t-test was used to find out the statistical significance between post and post-t-test values of VAS and WOMAC.

Data Presentation

RESULTS

The Paired 't' test analyses for the pre-test and post-test variable for the visual analogue scale for measuring pain in knee osteoarthritiss which is shown in table I. Both groups show a significant difference between the pre-test and post-test values. The 't' value for Group A is 14.the 't' value for Group B is 36.5. The unpaired 't' test analyses for the





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Post-test variables for Both groups for the visual analogue scale for measuring pain in the knee are shown in table 3. There was a significant difference shown between the Groups. Group B subjects show superior to Group A. The 't' value for the post-test variables for both groups is 7.4255. The Paired 't' test analyses for the pre-test and post-test variables for the disability measurement by WOMAC are shown in table 2. Both groups show a significant difference between the pre-test and post-test values. The 't' value for Group A is 21.12, and the 't' value for Group B is 52.62. The unpaired 't' test analyses for the Post-test variables for Both groups' disability measured by WOMAC is shown in table 3. There was a significant difference shown between the Groups. Group B subjects show superior to Group A. The 't' value for the post-test variables for both groups is 12.85.

DISCUSSION

The goal of this study was to determine how mulligan's mobilisation with movement and whole-body vibration treatment affected participants with knee osteoarthritis patients' levels of pain and impairment. Using the predefined inclusive and exclusive criteria, 30 participants were chosen. Ten each were assigned to two groups of subjects. Whole-body vibration exercises were performed by Group A, and Mulligan mobilisation with movement was performed by Group B. There was a substantial decrease in pain and disability between the pre-and post-treatment values in both Groups, which was validated by the following research. According to my research, there was a substantial decrease in pain and impairment in both groups when it came to whole-body vibration and mulligans mobilisation with movement, as seen by the post-test results. According to studies, WBV regulates neuromuscular responses, enhancing muscle strength. A probable explanation for the beneficial benefits of WBV on knee OA is the ability of the vibration created by the oscillating platform to stimulate muscle spindles, which activates the alphamotor neuron. This is followed by the vibration tonic reflex and spinal and supra spinal processes[10]. A subgroup analysis revealed that both high- and low-frequency WBV were successful in reducing pain, enhancing bodily function, and strengthening the knee extensor muscles. Numerous vibration characteristics may affect the efficiency of WBV on results, according to studies[11]. The study revealed that MWM combined with exercise improved functions more than other therapies did. The fact that MWM treatment for osteoarthritis significantly reduced pain and stiffness and enabled the subjects to move more freely without experiencing pain is one of the key factors contributing to the improvement in functional status. This allows the subjects to carry out exercises and other daily activities more successfully and pain-free[12]. The outcome of this study's data analysis shows that Mulligan's mobilisation with movement has a better impact than whole-body vibration. The substantial improvement is therefore founded on his idea. This study found that Mulligan mobilisation with movement was more effective at reducing pain and impairment than whole-body vibration workouts.

CONCLUSION

This study compared the effectiveness of Mulligan's mobilisation with movement and whole-body vibration treatment in reducing pain and impairment in participants with knee osteoarthritis. 30 participants between the ages of 45 and 55 were chosen following a careful examination of the inclusion and exclusion criteria. Two groups were created out of the subjects. Group B received mulligan mobilisation with movement whereas Group A received whole-body vibration. A VAS was used to quantify the level of pain. The WOMAC scale was used to assess functional abilities. Before the start of the treatment regime and after it had ended, the values of the outcome measures were noted. The data analysis appears to support the usefulness of mulligan's movement-based mobilisation and whole-body vibration in treating pain and functional limitations in people with knee discomfort. The fact that there has been a noticeable improvement in the groups demonstrating the usefulness of whole-body vibration and mulligan's mobilisation with movement in boosting functional activities and also reducing pain is evidence of this. Thus, it can be concluded that mulligan's mobilisation with movement and whole-body vibration are extremely efficient in treating patients with knee pain and lead to a considerable improvement in functional activities and pain relief.





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Table 1 Data Analysis and Presentation

Data values	VISUAL ANALOG SCALE (VAS)
Mean Difference	4.20
Standard Deviation	0.77
The unpaired t test	7.42
Table value	2.15

Table 2 Data Analysis and Presentation

Tubic 2 Dutu minury	old until I recentation
Data Analysis	THE WESTERN ONTARIO AND MCMASTER UNIVERSITIES OSTEOARTHRITIS INDEX (WOMAC)
Mean Difference	59
Standard Deviation	4
The unpaired t test	12.85
Table value	2.15





RESEARCH ARTICLE

A Multivariate Regression Analysis of RMT

G. Gomathi Eswari^{1*} and A. Rameshkumar²

- ¹Assistant Professor, Department of Mathematics, Srimad Andavan Arts and Science College, (Affiliated to Bharathidasan University) Tiruchirappalli, Tamil Nadu, India.
- ²AssistantProfessor, Department of Mathematics, Marudupandiyar College, Thanjavur (Affiliated to Bharathidasan University, Tiruchirappalli), Tamil Nadu, India.

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*Address for Correspondence

G. Gomathi Eswari

Assistant Professor, Department of Mathematics, Srimad Andavan Arts and Science College, (Affiliated to Bharathidasan University) Tiruchirappalli, Tamil Nadu, India. Email: mathseswari@gmail.com



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ABSTRACT

This paper presents investigations of the spectrum characteristics of random matrices, with a special emphasis on their most extreme eigenvalues, for both realistic and difficult examples of finite and infinite sample sizes, for a single Wishart matrix and for two Wishart matrices, respectively. The Tracy-Widom Laws, which characterise the distribution of the likelihood of the largest eigenvalue of a random matrix after normalisation, are intriguing to many individuals in RMT. By limiting the probability distributions of the eigenvalues of a certain random matrix, Wigner's semicircle law is produced.

Keywords: Spiled covariance model, Tracy-widom laws, Wigner matrix, Wigner's

semicircle law, Wishart matrix. MSC code: 15B52, 62H10

INTRODUCTION

We describe the spectrum of random matrices as the set of all eigenvalues, not just the ones in the middle. We set up a deterministic Markov chain to study the majority of the eigenvalues of the (n x n) Wishart matrix S = xx, which is constructed from a (n x r)-matrix X with entries that are iid standard Gaussian deviations (where r is the resolution and n is the sample size). Particularly notable are the outcomes for low n (exact distribution) and high n (asymptotic allocation including Wigner's semicircle rule). The eigenvalues of a single Wishart matrix can





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be analysed using multivariate statistics to characterise the distribution of their total values. Both fixed and large r scenarios as well as finite and enormous n cases are explored. The spiking covariance model of Iain Johnstone, which has lately become a hot topic in RMT, was also investigated. Then, for fixed r and limited sample sizes, we explain the situation of two independent Wishart matrices and their uses in multivariate regression and MANOVA. The spectral extrema for a single Wishart matrix and for two Wishart matrices are analysed, along with the largest eigenvalue distribution in a Gaussian ensemble and a Wishart-Laguerre ensemble. The Tracy-Widom Laws are valid for very large r and n. Computational programs for calculating the distributional findings of this study are discussed, and a quick overview of the RMT literature is provided.

Spectrum of Random Matrices

For large random matrices, the random behaviour of the spectrum's edges and centre warrants particular attention. A matrix's greatest and lowest eigenvalues are interesting as extremes, whereas the bulk of the eigenvalues is interesting. Whether we take into account the Gaussian orthogonal Ensemble (GOE), the Gaussian unitary Ensemble (GUE), or the Gaussian symplectic Ensemble, all the eigenvalues of Hn are real and may be ranked ordered. A square matrix's stability and invertibility strongly rely on its extremes. There is a difference between the statistics of the extreme eigenvalues in the spectrum's perimeter and the statistics of the spectrum's center, according to recent study.

The likelihood that consecutive eigenvalues are near together is low and the probability quickly declines as a power of the distance between them, which is an important characteristic that all three random matrix ensembles possess. Any two (correlated) eigenvalues extracted from a GOE (or GUE or GSE) matrix will exhibit this behaviour. While repulsion for iid points on an interval has a slower probability convergence rate to zero than repulsion for eigenvalues, both hold true. Therefore, spacings close to 0 are impossible given the distribution of spacings. Some features of quantum chaos are linked to this attribute.

Bulk of the Spectrum

The Real Wigner Matrix

The first focus of Wigner's study was the real symmetric (n x n)-matrix Hn = (Hij), in which diagonal entries are always equal to 0 and off-diagonal elements are always equal to 1 with probability 1/2. It was later deduced that such generalizations of these findings for this matrix would hold.

Gaussian ensembles as categorized by Dyson. The parameters $\beta \in \{1, 2, 4\}$ are used to classify the time-reversal invariance (TRI) and spin-rotational symmetry of the Hermitian matrix Hn = (Hij) and its matrix of eigenvectors U. (SRS)."Not appropriate" is signified by the word NA. Since then, a number of other definitions of a genuine Wigner matrix have been published. When we declare that "Hn is a Wigner matrix," we indicate that Hn belongs to the set of uncorrelated symmetric random matrices known as the Wigner ensemble. The entries on the diagonal of such symmetric matrices are always generated at random from the same distribution as the ones on the off-diagonal, but the off-diagonal entries may be drawn at random from any distribution. If the real wigner matrix is a member of the GOE, then it is a specific kind of real wigner matrix defined as a symmetric (i.e., Hn = H_n^T) random (n x n)-matrix where the ijth element is real-valued with distribution, $H_{ij}^{iid}N$ (0, σ_{ij}^2), $\sigma_{ij}^2 = 1 + \delta_{ij}$ where $\delta_{ij} = 1$ if i = j and otherwise. If a complex Wigner matrix belongs to the GUE, it is defined in a similar fashion. H_n^T

Finiten: Exact Distribution

Let U be the matrix of related eigenvectors and let $\lambda 1 > \lambda 2 > ... > \lambda n$ be the ordered eigenvalues of a real Wigner matrix Hn. Then, Hn = U Λ U τ , where Λ = diag{ $\lambda 1$, . . . , λn }. We can see that the sole factor that influences $tr[V(Hn)] = \sum_{j=1}^{n} V(\lambda j)$ is the eigenvalue. The assumption that the eigenvectors are uniformly distributed among the





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elements of each matrix ensemble has no bearing on the distribution p(Hn). The accurate joint probability distribution of the eigenvalues is obtained by multiplying the Jacobian of the form matrix by p(Hn), which also yields the eigenvalues and eigenvectors of a $(n \times n)$ Wigner matrix. The eigenvalues are distributed exactly as follows,

P (
$$\lambda_1,...,\lambda_n$$
) = $c_n e^{-1/4} \operatorname{tr}\{H_n\} \prod_{1 \le j < k \le n} |\lambda_j - \lambda_k|$, (1)
Where
 $\operatorname{tr}\{H_n^2\} = \sum_{j=1}^n \lambda_j^2$ (2)

and the normalizing constant, c_n , is dependent upon n.

For generic β -Gaussian-Hermite ensembles with β = 1 (GOE), 2 (GUE), or 4 (GSE), the Dyson index is supplied by the joint probability density of the eigenvalues of Hn.For generic β -Gaussian-Hermite ensembles with β = 1 (GOE), 2 (GUE), or 4 (GSE), the Dyson index is supplied by the joint probability density of the eigenvalues of Hn.

$$p_{\beta}(\,\lambda_1,\ldots,\lambda_n) = c_n \beta e^{-\frac{\beta}{4}} \mathrm{tr}\,\,(H_n^2) \ \prod_{1 \leq j < k \leq n} \ |\lambda_j - \lambda_k|^{\beta} \quad (\,3\,)$$

Where the normalizing constant,

$$c_{n,\beta} = \frac{1}{2\pi^{n/2\beta^{n}/2+\beta^{n}(n-1)/4}} \prod_{i=1}^{n} \frac{r(1+\frac{8}{2})}{r(1+\frac{8i}{2})}$$
 (4)

is dependent upon n and β .Dyson's β ensemble is another name for the distribution in (3). In certain discussions of this topic, the term is replaced with = $2/\beta$, leading to the outcomes for GOE α = 2, GUE α = 1, and GSE α = 1/2.

Large n: Wigner's Semicircle Law

We will examine the distribution of the empirical eigenvalues of a real (nn) Wigner matrix, n-1/2Hn, as n draws nearer to infinity. Let IA be the function that indicates that event A has occurred, and let $\#\{\cdot\}$ be the total number of items in the set being indicated.

The ESD of n-1/2Hn, $Gn(\lambda)$, converges asymptotically to the non-random limiting distribution $G(\lambda)$, according to Wigner's conclusion.

$$G_{n}(\lambda) = \frac{1}{n} \#\{i : \lambda_{i} \leq \lambda\} = \frac{1}{n} \sum_{i=1}^{n} I_{[\lambda i \leq \lambda]} \xrightarrow{a.s} G(\lambda), n \to \infty, \tag{5}$$

Where G(k) has density

$$g(\lambda) = \frac{1}{2\pi} \sqrt{4 - \lambda^2}, |\lambda| \le 2.$$
 (6)

and zero for $|\lambda| > 2$. This limiting density, which is a semicircle of radius 2, is known as Wigner's semicircle law. In free probability theory, Wigner's semicircle rule may be seen as the standard Gaussian's matrix counterpart.





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The kth Catalan number is the 2k-th moment of $g(\lambda)$,

$$C_K = \frac{1}{K+1} {2K \choose K} \tag{7}$$

and, by symmetry, the 2k + 1-st moment is zero, k = 0,1,2,..., where $C_0 = 1$ by convention. Numbers 1, 1, 2, 5, 14, 42, 132,... are called Catalan numbers because they follow the recursion formula $C_{k+1} = \sum_{i=1}^{K} C_{i-1} C_{k-i}$. It turns out that the only distribution for which the Catalan numbers serve as even moments is Wigner's semicircle law.

The Method of Moments: This was Wigner's first evidence (1955). Let $gn(\lambda)$ be n1/2Hn's empirical spectral density function. This method takes use of the ability of expressing the kth instant of $gn(\lambda)$ as

$$\int_{-2}^{2} \lambda^{k} g_{n}(\lambda) d\lambda = \frac{1}{n} \operatorname{tr} \left\{ (n^{-1/2} H_{n})^{k} \right\} = \frac{1}{n^{1+k/2}} \operatorname{tr} \left\{ H_{n}^{k} \right\}.$$

As $n \to \infty$ starts going down, the moments of $gn(\lambda)$, which were given above as the normalised trace of powers of Hn, converge to the Catalan numerals, which are the even moments of the semicircle law $g(\lambda)$.

The Stieltjes Transform Method: The finite-n Stieltjes transform of n1/2 Hn can be written as

$$S_n\left(z\right) = \int_{-\infty}^{\infty} \frac{g \mathbf{n}(\lambda)}{\lambda - z} \, \mathrm{d}\lambda \ = \frac{1}{n} \, \mathrm{tr} \, \{ \left(n^{-1/2} H_n - z I_n \right)^{-1} \},$$

For a simple (nn) Wigner matrix, convergence to the semicircle law is seen in Figure 1 above. To determine the eigenvalues of Hn, we first sampled n = 1,000 iid standard Gaussian deviations (left panel) and n = 25,000 (right panel). For both scenarios, we provide histograms of the Hn eigenvalues.

Large n: Largest Eigenvalue

the circumstances that must be met for the highest eigenvalue $\lambda 1$ of the normalised Wigner matrix Hn(n-1/2) to converge to a constant as $n \to \infty$. Mean zero and a finite fourth moment for entries off the diagonal and a finite second moment for entries on the diagonal were prerequisites.

Single Wishart Matrix

A random r-vector X is of relevance in multivariate statistics if it follows a distribution with a mean vector μ and a covariance matrix Σ of the form where

$$\mu = E\{X\}, \ \Sigma = E\{(X-\mu)(X-\mu)\tau\}.$$
 (8)

That X must also be Gaussian is an assumption. Numerous applications of multivariate analysis make use of dimensionality reduction and the study of functions of Σ , including its eigenvalues and associated eigenvectors.

Usually, Σ must be calculated using a sample of data since it is unknown. The standard estimate of is given by where X is an independent random variable chosen from the same distribution as the r-vectors Xi,i = 1, 2,..., n.

$$\widehat{\Sigma} = n^{-1} \sum_{i=1}^{n} (X_i - \bar{X}) (X_i - \bar{X})^{\mathrm{T}} = n^{-1} X_c X_c^T,$$
(9)

Where $\bar{X} = n^{-1} \sum_{i=1}^{n} X_i$ is the sample mean vector and is the population mean vector that may be estimated using the sample mean. $X = (X_1, \dots, X_n)$ and $X_0 = X(I_n - n - 1J_n)$, where $J_n = I_n I_n \cap T$ and I_n is an n-vector of I_n . In order to examine how the eigenvalues of I_n change under various assumptions for the number of variables I_n and the number of observations I_n , we may convert I_n into an unbiased estimator of by replacing I_n with I_n

A Spiked Covariance Matrix

As a specific alternative to the "null" model of a white Wishart distribution, where = Ir, the "non-null" idea of a "spiked" covariance model is presented. The bulk of the eigenvalues of the covariance matrix in this model are 1. The





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spiking covariance model has shown promise in a variety of fields, including mathematical finance, wireless communication, signal processing, and speech recognition.

Assume m, the number of spikes, is known and that the first m eigenvalues are unique. Let Jm = $\{\lambda 1, \dots, \lambda m\}$. In the null model, $\lambda 1 = \lambda 2 = \dots = \lambda m = 1$. The covariance matrix, m, of a population that is not null is a change to the identity matrix with a finite number of rows. One alternative to using the spiking covariance model to characterise the non-null case is to rescale it. The lowest eigenvalues ($\lambda m+1,...,\lambda r$) don't take the value 1; instead, they take the value $\sigma 2$.

Two Wishart Matrices

Fixed r, Finite n: Exact Distribution

Real case. Let X_1^{iid} $N_r(0,\Sigma)$, i=1,2,...,n, and let $X=(X_1,\cdots,X_n)$ be an $(r\times n)$ -matrix. Let $A=XX^\tau \sim W_r(n,\Sigma)$. we have another $(r\times r)$ -matrix $B \sim W_r(m,\Sigma)$ that is independent of A. We may select $\Sigma = Ir$ without sacrificing generality if our focus is on the eigenvalues of A-1B, whose distribution does not rely on.

We now have two white Wishart matrices, $A \sim Wr(n,Ir)$ and $B \sim Wr(m,Ir)$, where A, B, and their product A+B are invertible if and only if m, $n \ge r$. The following generalized eigenvalue issue (for λ) is of relevance to ourselves:

$$|B-\lambda(A+B)| = 0.$$
 (10)

The eigenvalues of the matrix (A + B)-1B are of importance to us. It follows that $0 < \lambda < 1$ in (10) because A is positive definite. This is a different way to represent the eigenequation (10):

$$|B - \theta A| = 0,(11)$$

and, in this form, we are interested in the eigenvalues of $A^{-1}B$ The eigenvalues λ and θ are related by $\lambda = \theta/(1+\theta)$ or $\theta = \lambda/(1-\lambda)$.

The eigenvalues of the generalised eigenequation (10) are exactly distributed in a joint manner by

$$p(\lambda_1,...,\lambda_r) = c_{m,n,r} \prod_i \left[w_{a,b} (\lambda_i) \right]^{\frac{1}{2}} \prod_{i < j} |\lambda_i - \lambda_j|, \tag{12}$$

where

$$w_{a,b}(\lambda) = \lambda^a (1-\lambda)^b, a = n - r - 1, b = m - r - 1,$$
 (13)

is a normalising constant that relies on m, n, and r, and is a weight function for the Jacobi family of orthogonal polynomials.

For a more general form of (12), we get

$$P\beta(\lambda_1,...,\lambda_r) = C_{m,n,r,\beta} \prod_i \left[w_{r,m,n,\beta}(\lambda_i) \right]^{1/2} \prod_{i < j} |\lambda_i - \lambda_j|^{\beta}, \quad (14)$$

Where

$$W_{r,m,n,\beta}(\lambda) = \lambda^a (1-\lambda)^b$$
, $a = \beta(n-r+1)-2$, $b = \beta(m-r+1)-2$, (15)

and $c_{m,n,r,\beta}$ is a normalizing constant that depends upon m, n, r, and β . The values of are the same as before: β = 1 for reals, β =2 for complexes, and β = 4 for equations.

Setting $\lambda_i = \theta_i/(1 + \theta_i)$, the joint distribution of the eigenvalues of the generalized eigenequation (12) is given by

$$p(\theta_1,...,\theta_r) = c_{m,n,r} \prod_i [w_{a,b}(\theta_i)]^{1/2} \prod_{i < j} |\theta_i - \theta_j|,$$
 (16)

where





(17)

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$$W_{a,b}(\theta) = \theta^a(1+\theta)^b$$
, $a = n-r-1$, $b = m+n$.

By substituting $\theta = (1 + x)/2$ for in Equations (16) and (17), we get the Jacobi orthogonal ensemble.

$$p(x_1,...,x_r) = c_{m,n,r} \prod_i [w_{a,b} (x_i)]^{1/2} \prod_{i < j} |x_i - x_j|,$$
 (18)

where

$$wa,b(x) = (1-x)a(1+x)b, a = n-r-1, b = m-r-1,$$
 (19)

is the weight function for the Jacobi family of orthogonal polynomials.

Applications

Example: 4.1

The Patience sorting algorithm was developed from a British card game played by a single player. John Hammersley, seeing the potential in this, created the ensuing method to determine how long the largest rising subsequence is. Subadditive ergodic theory was also inspired by this finding.

The initial situation involves a "deck" of cards bearing the digits 1, 2, 3,..., n. After shuffling the cards, one is drawn at random and placed in a pile according to the rules below. Only cards with a value lower than the card on top can be added to a pile that already contains face-down cards. In the event that it isn't, a new pile is created to the right of the existing ones. It is possible to stack threes on top of fives, but a six requires you to start a new stack. Until all of the cards have been dealt and piled, the game will continue. In this game, you have to get rid of as many piles as possible.

Example: 4.2

Assume the previous game's order of cards was 2, 5, 1, 3, 4, 8, 6, 7. The 2 should be used to initiate the first pile; the 5 is larger than the 2, so it should initiate the second pile; the 1 should be placed atop the 2 and the 3 should be placed atop the 5; the 4 should initiate the third pile; the 8 should initiate the fourth pile; the 6 should be placed atop the 8; and the 7 should initiate the final pile. The piles are as follows:

The best approach is the "greedy" one, in which each card is placed on the leftmost pile possible. As with the longest rising subsequence of the previous permutation, the outcome in this case is five heaps. In most cases, the optimum greedy technique will result in a number of piles that is equal to the longest rising subsequence.

Example: 4.3

Cards may not be stacked in ascending subsequence order at the top of their respective heaps if they are not in permutation order. Here's a look at the numbers 8, 6, 1, 3, 4, 7, 5, 2, 8, and 6. The piles are:





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It's simple to see that the numbers bolded (1, 2, 4, 5) do not form a continuous subsequence of the main sequence, which progresses from least to largest. But the lengths of the subsequences 1, 3, 4, 5 or 1, 3, 4, 7 also equal 4, hence the number of piles is also 4. It turns out that there is a highly fast technique for determining the length of the longest rising subsequence, and that this corresponds to patience sorting $L_n = l_n(\pi)$.

Example: 4.4

Distances between the Riemann zeta function's successive zeros. By summing inverse powers of the positive integers, Riemann's zeta function is defined.

$$\zeta(s) = \sum_{n=1}^{\infty} n^{-s} = 1 + \frac{1}{2^8} + \frac{1}{3^8} + \frac{1}{4^8} + \dots$$
 (20)

Set s = 2, and Leonhard Euler's famous finding that $\zeta(2) = \pi 2/6$ holds. The zeta function, as Euler showed, may also be represented as a product over the primes.

$$\zeta(s) = \prod_{P} (1 - p^{-s})^{-1} = \frac{1}{\left(1 - \frac{1}{2^{8}}\right)\left(1 - \frac{1}{3^{8}}\right)\left(1 - \frac{1}{5^{8}}\right)\left(1 - \frac{1}{7^{8}}\right)\dots}}$$
(21)

The product of Euler is a well-known formula. As a result, the zeta function has characteristics with the pattern of prime numbers. But the storey has more to it. Since the zeta function has zeros on the complex plane, Riemann proved that it may be expressed as a product over those zeros.

$$\zeta(s) = f(s) \left(1 - \frac{8}{p^1}\right) \left(1 - \frac{8}{p^2}\right) \left(1 - \frac{8}{p^3}\right) \dots$$
 (22)

where $\varrho 1$, $\varrho 2$,... are the complex integers for which $\zeta(s) = 0$, and f(s) is a straightforward "fudge factor."

CONCLUSION

We calculate the eigenvalues of a white Wishart matrix, the Markov-Pastur semicircle laws, the Tracy-Widom distributions, and Wishart-matrix simulations. Despite the book's focus on Gaussian ensembles, both Laguerre ensembles and Wishart matrices are mentioned for the crucial roles they play in mathematical statistics. All the best papers ever written on RMT are collected here.

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TABLE1

β	Ensemble	TRI	SRS	H_{ij}	U
1	GOE	Yes	Yes	real	orthogonal
2	GUE	No	NA	complex	unitary
4	GSE	Yes	No	real-quaternion	symplectic

Table 2 Families of orthogonal polynomials and their weight functions w(x).

Case	w(x)	Interval	OrthoPoly
Gaussian	e-x2	$(-\infty,\infty)$	Hermite
Wishart	$x^a e^{-x}$	$[0,\infty)$	Laguerre
Two Wisharts	xa(1-x)b	(0,1)	Jacobi





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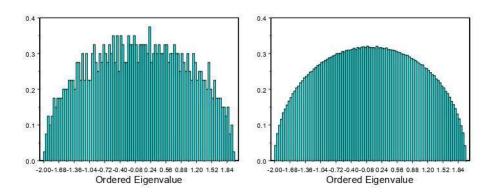


Fig. 1. Illustration of the convergence to Wigner's Semicircle Law. Normalized histograms of the eigenvalues from a single $(n \times n)$ Wigner matrix. Left panel: n = 1,000. Right panel: n = 25,000. For each n, there are 100 bins.

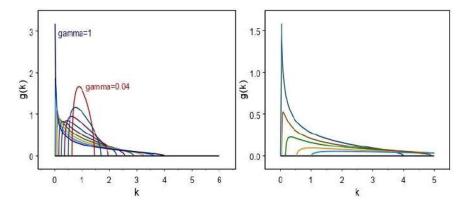


Fig. 2. Density of eigenvalues from the Mar cenko-Pastur Law. Left panel: $\gamma = 0.04, 0.09, 0.16, 0.25, 0.36, 0.49, 0.64, 0.81, 1 \ (i.e., r \le n)$. Right panel: $\gamma = 1, 1.5, 2, 3, 4 \ (i.e., r \ge n)$.

The spiked covariance matrix is given by $\Sigma_m = \text{diag}\{\lambda_1,...,\lambda_m,1....,1\}, \lambda_1 > \lambda_2 > \cdots > \lambda_m > 1$,





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REVIEW ARTICLE

Comprehensive Review: to Enhance Absorption and Bioavailability of Herbal Drug by Topical and Oral Route using Phytosome

Manisha Katkade^{1*}, Rahul shivarkar² and Shashikant Dhole³

Student, Department of Quality Assurance, PES Modern College of Pharmacy, (for Ladies) (Affiliated to University of Pune) Pune, Maharashtra, India.

²Assistant Professor, Department of Pharmacognosy, PES Modern College of Pharmacy, (for Ladies) (Affiliated to University of Pune) Pune, Maharashtra, India.

³Principal, Department of Pharmaceutics, PES Modern College of Pharmacy, (for Ladies) (Affiliated to University of Pune) Pune, Maharashtra, India.

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*Address for Correspondence Manisha Katkade

Student,

Department of Quality Assurance,

PES Modern College of Pharmacy, (for Ladies) (Affiliated to University of Pune)

Pune, Maharashtra, India.

Email: manishakatkade0704@gmail.com



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ABSTRACT

Phytopharmaceuticals or herbal drugs have a significant therapeutic effect on healthcare System. But due to poor lipophilicity and high polarity the active constituents are poorly absorbed resulting in poor bioavailability. These problems can be over comed by formulating a suitable novel preparation of the herbal extract i.e., Phytosome. Phytosome are one of the novel drug delivery system containing hydrophilic bioactive phyto constituents of herbs which are surround and bound by phospholipids. Phytosome technology is promising targeting novel drug delivery with improved efficacy, quality and target ability of active plant constituents. Novel herbal formulation techniques have assured the researchers to deliver the plant based secondary metabolites to their systemic targets for specific effect. The recent development and conducted works of various researchers have been studied thoroughly to establish the transdermal route as a potential way to deliver phytoconstituents to their specific targets. Phytosome can before emulated both topically and orally. Phytosome given by oral route increases the bioavailabilty of herbal drug. This review highlights the mechanism of action of phytosome by topical and oral route and also highlights unique properties of phyto-phospholipid complex along with their application in the novel natural drug delivery.

Keywords: transdermal, formulation, phytosome, therapeutic, plant





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INTRODUCTION

The capability of the dosage form to deliver the medicament to its specific site of action at a rate and amount sufficient to achieve the desired curative effects is critical to the efficacy of any drug, whether it is comes from a plant, animal, or synthetic drug. There are numerous innovative drug delivery systems and methods can be supplied through various channels to allow the medication to pass through the barrier at appropriate and safe dosages. Phytopharmaceuticals or herbal extracts(Drugs) have a significant therapeutic effect on healthcare systems. The herbal extracts and its active constituents show excellent pharmacological in vitro effects, they still have very poor in vivo biological effects because of their considerable molecular weight and low lipid solubility, it leads to low systemic availability[1]. Additionally, combination of plant extracts into delivery channels will help improve and bulk dosing of these phytochemicals, as the activity of the herbal medicines depends on the overall synergism effect of all the constituents. Hence, integrating innovative delivery systems with traditional medicines is very important, especially for treating chronic diseases like asthma, diabetes, osteoporosis, and age-related diseases like dementia[2]. Novel drug delivery system is a novel approach to drug delivery which addresses the limitations of the traditional drug delivery systems. Phytosome is a novel patented technology which is developed to incorporate standardized plant extracts into phospholipids to form a lipid soluble molecular complexes, with improved bioavailability and absorption, which is called as "phytosome". The term "Phyto" means plant, and "some" i.e., cell-like. Phytosome formed from the reaction of stoichiometric amount of phospholipid with standardized herbal extract or polyphenolic constituents in an aprotic solvent. Phytosome shows better stability due to formation of chemical bonds between phosphatidylcholine molecules and phytoconstituents[3]. The phytosome technology was first developed by the Indena S.p. A, Italy. The polyphenols active pharmaceutical ingredients are slightly soluble in both water and lipids.

This phenomenon makes the polyphenolic APIs difficult to be formulated into commercial medicines. In 1991, Bombardelli and Spelta have formulated a new drug delivery system and named it phytosome[4]. A phytophospholipid complex (Phytosome) is the promising emerging technique applied to pharmaceuticals for the enhancement of bioavailability of herbal extracts and phyto constituents. Phytosome are the advanced forms of herbal formulations contains the bioactive phyto constituents of herbal extract and it has good ability to transition from a hydrophilic into the lipophilic environment of the cell membrane, it exhibit better pharmacokinetic and pharmacodynamic profile than conventional herbal extracts. Phytosome formulated in the form of various dosage forms like suspensions, capsules, creams, gels etc[5]. The Phytosome increases the absorption of active ingredients of herbal extract, when topically applied on the skin, And it also improves systemic bioavailability when administered orally[5]. Nanoparticles and Nanomaterials are Significant advancements have gained attention for the design of delivery methods because of their superior biological and chemical performance. Nano carriers includes safe and inert materials investigated for their potential role in medication(Drug) delivery and unique characteristics. A nano liposomal delivery system i.e.,Phytosome. is one such system for improved site-specific delivery of phytochemicals with enhanced absorption for the topical and oral route[6]. The updated pharmacokinetic and therapeutic parameters of phytosome are becoming beneficial in medical and cosmetic industries[8].

STRUCTURE OF PHYTOSOME

The structure of phytosome is majorly composed of standardized polyphenolic plant extract incorporated into phospholipids, mainly phosphatidylcholine(PC)[8]. The phytosome in the form of lipid vesicles are the result of a H-bond interaction between the polyphenolic moiety of the bioactive herbal extracts and the phosphate group of phospholipids matrix in non-polar solvents[9]. The water-soluble polyphenolic rings of phytochemicals (i.e., flavonoids and terpenoids) having a high affinity to chemically bind towards to the hydrophilic moiety of phospholipids (i.e., choline) to form the body of phytosome, while the phosphatidyl lipophilic moiety of the phospholipids forms a tail to incorporate the water-soluble choline bound phytoconstituents[10-11]. The encapsulated form(Capsule) of poorly soluble polyphenolic compounds into the phytosomal delivery system has a significant effect on the enhancement of their absorption, and it leading to better penetration and absorption across the biological membrane and enhanced bioavailability [12].





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The potential role of Phytosome is in the improvement of herbal-originated polyphenolic compounds used for the treatment of several diseases and makes this nanotechnology a promising tool for the development of new formulations[13].

COMPONENTS

Phospholipids

Phospholipids which are extracted from soy, mainly phosphatidylcholine(PC), are lipophilic substances and readily complex polyphenolics. In this context, phosphatidylcholine(PC) are the major molecular building block of cell membranes and it is miscible in both water and in oil/lipid environments, and is well absorbed orally, and it has the potential to act as a chaperon for polyphenolics, accompanying them through biological membranes[14]. Phospholipids are majorly present in egg yolk and plant seeds. Phospholipids can be divided into glycerophospholipids and sphingomyelins depending on the backbone. Additionally, glycerophospholipids include phosphatidylcholine(PC), phos-phatidylethanolamine(PE), phosphatidylserine (PS), phosphatidyl- glycerol (PG) and phosphatidic acid (PA)[15] . PC, PS, and PE are the major phospholipids used to prepare complexes which are composed of a hydrophilic head group and two hydrophobic hydrocarbon chains[16]. Among these three phospholipids, PC is the most frequently used to prepare phospholipid complexes(Phytosome). The benefits of PC include it shows amphipathic properties that give it moderate solubility in water and lipid media. Also PC is an essential component of cell membranes, and accordingly it also exhibits robust biocompatibility and low toxicity[17].

Phyto-active constituents

The active constituents of herbal extracts identified by the researchers are generally defined based on robust *in vitro* pharmacological effects, rather than on *in vivo* activities. Most of these phyto active constituents are polyphenols. Some of the biologically active polyphenolic constituents of plants shows affinity towards the aqueous phase and cannot pass through biological membranes, such as hesperidin. And others have high lipophilic properties and cannot dissolve in aqueous gastrointestinal fluids, such as curcumin and rutin[17].

Solvents

Different solvents have been utilized by different researchers as the reaction medium for formulation of phytophospholipid complexes(Phytosome). Traditionally, aprotic solvents, such as aromatic hydrocarbons, methylene chloride, halogen derivatives, ethyl acetate, or cyclic ethers etc. have been used to prepare Phytosome but they have been largely replaced by protic solvents like ethanol[18-19]. Protonic solvents, for e.g., ethanol and methanol, have been more recently been successfully utilized to prepare phospholipid complexes. For example, the Xiao prepared silybin-phytosome using ethanol as a protonic solvent; subsequently, the protonic solvent was removed under vacuum at 40 °C[20].

Stoichiometric ratio of active constituents and phospholipids

Normally, Phytosome are employed by reacting a synthetic or natural phospholipid with the active constituents in a molar ratio ranging from 0.5 to 2.0 [21]. Whereas, a stoichiometric ratio of 1:1 is considered as the most efficient ratio for preparing phospholipid complexes[22]. For example, quercetin-phospholipid complexes(Phytosome) were prepared by mixing Lipoid S 100 and quercetin at a molar ratio of 1:1[23]. However, different stoichiometric ratios of active constituents and phospholipids have been used to prepare different Phytosome.

DIFFERENT METHODS OF PREPARATION

The methods used for phytosome preparation are described by the, Jiang *et al* [24], Maiti *et al* [25] and Maiti *et al* [26]. In the 2006, Yanyu *et al* prepared a silybin-phospholipid complex using ethanol as a reaction medium. Silybin and phospholipids were mixed into the medium, after the organic solvent was removed under vacuum condition, and a silybin Phytosome was formed[27]. Common stages in formulation of Phytosome are described in following Figure 1 Various methods of preparation of phytosome are as follows:-





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Anti-solvent precipitation process

Take Specific amount of herbal extract and phospholipids and refluxed it with 20 ml of organic solvents such as acetone at specific experimental conditions below 50°C for 2-3 hrs. Then concentrate reaction mixture to minimum volume up to 10 ml and then add solvent with low polarity such as n-hexane with stirring, precipitates are obtained and filter it. Filtered precipitates are stored in desiccators or freezer. The dried precipitates are grinded and powdered complex are stored in dark amber colored glass bottle at room temperature [28].

Rotary evaporation process

Take Specific weight of herbal extract and phospholipids were mixed in 30 ml water miscible organic solvent such as acetone in round bottom glass container and stir it for 2 hours at a temperature less than 50°C in rota evaporator. Anti solvent such as n-hexane added to thin film which is obtained after uninterrupted stirring using a stirrer[29]. Precipitate of phytosome which obtained can be stored in amber colored glass container at controlled temperature under specified humidity.

Solvent ether-injection process

This process involves reaction of lipids dissolved in organic solvent with herbal extracts in aqueous phase. Phospholipids are solubilised in diethyl ether and are slowly injected drop wise in an aqueous solution of the phytoconstituents which is to be encapsulated to form phytosome. It results in the formation of cellular vesicles on subsequent solvent removal, leading to complex formation(Phytosome)[30]. Structure of phytosome depends upon concentration, when the concentration is less amphiphiles in mono state are produced, but variety of structures with different shapes *i.e.*, cylindrical, disc, round and cubic or hexagonal vesicles may be formed on increasing the concentration [31].

NOVEL METHODS

Novel methods for the phospholipid complexation which include supercritical fluids which include gas solvent technique, compressed solvent process and supercritical solvent method[31]. Traditional(Conventional) methods have several drawbacks, such as its including multistep processes, difficulty extraction, and time consumption. Supercritical fluid methods are used to change the size, morphology and shape of material of interest. Along with other benefits like as a single-step process, the capacity to process thermolabile substances, and eco-friendly technology, high product purity, crystal polymorphism controls Supercritical fluid (typically CO2) used as an antisolvent in the Gas anti-solvents technique (GAS), Supercritical antisolvent technique (SAS), and Solution enhanced dispersion by supercritical fluids (SEDS) techniques, also Rapid expansion of supercritical solutions (RESS) uses it as a solvent [32].

Gas anti-solvents approach (GAS)

Supercritical CO2 gas usage is not necessary when using CO2 as an antisolvent. To accomplish homogeneous mixing, it is injected into the solution in a closed chamber, preferably from the bottom. Solutes precipitate because of the decreased solubilization power of the organic solvent brought on by the dissolution of CO2 gas. To get rid of any remaining solvent, the particles are rinsed with more antisolvent. If not, the solutes could resolubilize during the depressurization step, endangering the stability of the product. When scaled up to industrial levels, the gas antisolvent technology outperforms the solvent antisolvent technique in terms of results[33].

Supercritical antisolvent precipitation (SAS)

Submicrometer-sized particles with a restricted size distribution are produced by removing the solvent from the gas phase by reducing the pressure in the SAS. The supercritical state of CO2 is required. The solution and CO2, both are injected from the top into a closed chamber. In contrast to GAS, this method has a proven record of widespread success[34].





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Mechanism of Action of Phytosome by Topical and Oral Route

One of the most significant advances in the study of transdermal drug delivery systems (TDDS) is the increased bioavailability of phytoconstituents through the transdermal route. Therapeutic agents are delivered via the skin to the systemic circulation in TDDS. Due to its lipophilicity, the stratum corneum, the outermost horny layer, presents the main obstacle to medication transport. The phytoconstituent's phytosomal form aids in avoiding this barrier feature. The following summarizes the fruitful research experience of approximately twenty years of study by several researchers and shows that the phytosome technology is potent enough to be applied transdermally to increase the bioavailability of phytoconstituents[35]. The skin is an excellent site for the administration of pharmaceutical agents or medication for both local and systemic impacts and it serves as a strong barrier to the permeability of the majority of substances[36] [Figure 2]. The transdermal permeability can be increased when the active ingredients are lipophilic and they have a low molecular weight. Bombardelli et al. noted in 1991 that phospholipids had a strong affinity for particular types of flavonoids. By complexing phospholipid with extremely polar plant derivatives—that is, complexes between a pure phospholipid and a pure active principle—they created a novel class of molecules known as phytosome. He first obtained some pharmacological information and established some chemico-physical characteristics of phytosome of escin, quercetin, catechin, and glycyrrhetinic acid. He stated right away that phytosome are an effective way to transport phytoconstituents over the skin[37]. Yanyu et al. (2006) looked at how, in silymarin phytosome in experimental models, the well-known calming effect of silymarin is enhanced by a factor of more than six. Because the complex has a stronger affinity for skin phospholipid[38], the phytosome form's activity is improved as compared to the free active principle. According to Kidd's (2009) findings, the phospholipid complex's liposomal-like characteristics are connected to the hydration of the superficial corneous layer. Because of their transdermic effect, ginselect phytosome facilitate the skin's absorption of the ginseng saponin found in the phospholipid complex[39] Phytosome offer a number of possible benefits over traditional topical formulations in topical treatments. Phytosome promote the transport of herbal active ingredients to tissues and improve skin absorption and bioavailability [40]. Additionally, phytosome strengthen collagen structure, the balance of enzymes, and moisture in the skin [41]. When compared to other free chemicals, the potency of phytosome was enhanced due to their strong affinity for skin phospholipids [42]. As was already indicated, topical uses of phytosome formulation are confronted with a number of challenges. For instance, the SC, or thick outer layer of the epidermis, is one of the most significant obstacles to the transdermal administration of phytochemicals [43]. There are two main routes that bioactive compounds might pass through the SC: intracellular or intercellular[Figure 3]. Sweat can facilitate the entry of cells into one another.

Sweat glands, sebaceous glands, or hair follicles can all be used for intercellular penetration, while cornecytes and the intercellular lipid matrix are the primary routes for intracellular penetration[44-45]. According to reports, increasing the drug's diffusion coefficient can boost biomolecule concentrations and improve the way these molecules are partitioned between the SC layer and the molecules themselves. Taken together, these factors can increase the biomolecule permeability to the SC for transdermal application [46]. Topical medicines with lipophilic and low molecular weight active components have the potential to enhance transdermal permeability. Due to their hydrophilic character, polyphenols, which make up the majority of the phytocompounds that have been extensively researched, have poor bioavailability and lipid solubility, which limits their in vivo activity[47]. Several flavonoid compounds can be strongly bound by the phospholipid moieties of phytosome with a high affinity [48]. Many herbal extracts, including ginseng, hawthorn, grape seed, green tea, milk thistle, and grape seed, work better when loaded into phytosome than when they are in a liposomal formulation. Because the phospholipid molecules in Phytosome interact with the active phytoconstituents, enhancing their stability, the formation of polyphenol-based phytochemicals phytosome nanoparticles improves the application of conventional herbal materials. Moreover, the phytosome-herbal combination has a greater affinity for the phospholipid moiety of the epidermis, hence enhancing the topical formulation's lipid solubility[49]. Topical dosage forms: A topical formulation of the Phytosome complex is also possible. Dispersing the phospholipidic complex in a tiny amount of the lipidic phase and adding it to the already-created emulsion at low temperatures (not higher than 40°C) is the optimal method for incorporating the Phytosome complex in emulsion. The primary lipidic solvents used in topical preparations are soluble in the





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Phytosome complexes. The Phytosome complex may also be disseminated into the watery phase and then added to the final formulation at a temperature lower than 40°C in formulations with a limited amount of lipids[50]. For topical use, phytosome have been developed as gel and cream According to Djekic et al., a hydrogel loaded with 18-GA phytosome was created by dispersing the phytosome and carbomer in water. 10% sodium hydroxide was then added to neutralize the formula. Hydrogel was formed later by adding humectant.

It is possible to manufacture Phytosome complexes for oral or topical use. To achieve the optimal results from this technological advancement in terms of formulation manageability and improved bioavailability (such as the proper disintegration and dissolving time of oral forms)

It has long been accepted that the best way to administer drugs is orally. Nowadays, there is continuous research being done on nano-formulations to improve absorption, increase bioavailability, and boost therapeutic efficacy. Formulation scientists have been drawn to lipid-based nano formulations because of their improved bioavailability and comparatively greater safety profile. The bioactives or medications can also be shielded from the urge of the gastrointestinal system by using these delivery methods. They also help hydrophobic medications that are trapped in lipid matrices to be absorbed. Lipid excipients have been shown to both increase the bioavailability of bioactives administered orally and decrease P-glycoprotein-mediated efflux[51]. Phytosome are advanced form which shows improvement in bioavailability of phytoconstituents. Polyphenolic phytoconstituents are complexed with phosphatidylcholine In 1:1 or 1:2 ratio, phytosome are formed. This is a novel patented technology of Indena, through which water soluble actives of plant origin can be delivered (mainly polyphenols). Most of the phytoconstituents (anthocyanidins,tannins, flavanoids and terpenoids) can penetrate lipoidal biological membranes due to their larger size and hydrophilicity. Phytosome are able to cross the hydrophilic and hydrophobic environments to successfully reach the human circulation system[51].

Phospholipid complex and their absorption by oral route

Phospholipid complexes are absorbed from the GIT through enterocyte based transport, and the drug transport to the systemic circulation via intestinal lymphatic system which has widespread network throughout the body. The major advantage of lymphatic transport i.e., to bypass the first-pass metabolism and applicable for targeted drug delivery. After oral administration of dosage form, the possible mechanism by which lipids affect drug bio availability is shown in Figure 4. Schematic diagram representing the possible mechanism by which phytophospholipid complex(Phytosome) entry into the intestine from unstirred water layer is both by direct solubilization through enterocytes or by endocytosis, inhibiting drug efflux by blocking transporter proteins, paracellular transport in lateral tight junctions, formation of chylomicron production and entering lymphatic port. Several researchers have proposed various processes for lipid absorption. Yeap et al. proposed that enterocytes play a role in lipid absorption. The paracellular transfer of phosphotidylcholine across lateral tight junctions was discovered by Stremmel et al.[51]. According to Holm et al., bile salts and other lipid emulsifiers can inhibit drug efflux by preventing the action of transporter proteins like CYP 450 and/or P-gp[52]. According to Jain et al., endocytosis facilitates medication absorption[53]. Green tea extract was formulated as Green Select Phytosome. and these were converted into oral coated tablets by Pierro et al. (2009). Clinical trial studies done on obese patients demonstrated that phytosome coated tablets were capable of weight loss when administered orally to patients [54]. In 2014. Keerthi B et al, Formulated and evaluated capsules of Ashwagandha Phytosome.

Oral Dosage Forms

Soft gelatin capsules

When creating Phytosome® complexes, soft gelatin capsules are a great option. To create solutions that may be put inside soft gelatin capsules, the Phytosome® complex can be distributed across greasy media. You can use semi-synthetic or vegetable oils for this. Granulometry of $100\% < 200 \mu m$ is recommended by Indena for optimal capsule manufacture. First feasibility trials should be carried out to determine the most appropriate vehicle because, based on Indena's experience, not all Phytosome complexes react in the same way when distributed in oily vehicles and when the oily suspension is packed in the soft gelatin capsules[51].





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Hard gelatin capsules

The Phytosome complex can be formulated in the form of hard gelatin capsules as well. Even though the phytosome complex's apparent low density appears to limit the amount of powder that can be placed into a capsule (typically no more than 300 mg for a size 0 capsule), a direct volumetric filling procedure (without pre compression) can be used. However, the amount of powder that can be put in a capsule using a piston tamp capsule filling method can be increased; however, pre compression may have an impact on the disintegration time. It is recommended by Indena to closely monitor the relevant parameters while developing new products and processes. To determine the optimal production technique, a preliminary dry granulation procedure isadvised[51].

Tablets

The best production method for producing tablets with greater unitary dosages and appropriate technological and biopharmaceutical characteristics is dry granulation. However, a direct compression process can only be used for low unitary doses due to the Phytosome complex limited flowability, potential stickiness, and low apparent density. It should be noted that whenever a direct compression process is used, the Phytosome complex should be diluted with 60–70% of excipients in order to maximize its technological properties and produce tablets with the proper technological and biopharmaceutical characteristics. However, because water and heat (granulation/drying) have a detrimental influence on the stability of the phospholipid complex, moist granulation should be avoided[51].

PROPERTIES OF PHYTOSOME

Hepatoprotective properties

The majority of PHYTOSOME® research focuses on the liver-protecting flavolignans found in Silybum marianum Gaertner. There are several flavonoids in milk thistle (Silybum marianum) fruit that have hepatoprotective properties [55-56]. Sily Research has demonstrated the efficacy of silymarin in the treatment of several kinds of liver conditions, such as cirrhosis, hepatitis, fatty infiltration of the liver (fatty liver caused by chemicals and alcohol), and bile duct inflammation [57-59]. Silymarin's antioxidant properties significantly improve the liver's tolerance to harmful incidents [60]. Based on three patterns of covalent interaction between its flavonol and lignan elements, silymarin has a complex composition.. Silymarin's primary and more potent component is silybin, which is a combination of two diastereomers [61]. Silybin maintains glutathione in the liver to protect It. In numerous human disease models, silybin has been found to be effective; however, due to its extremely poor oral bioavailability, the clinical application of these results has been prevented. In a pharmacokinetic study [62], the oral bioavailability of silybin was significantly increased when administered as silybin–phospholipid complex. The increased lipophilicity of the silybin–phospholipid complex has been claimed to explain the improved bioavailability. Silybin protects the liver by conserving glutathione in the parenchymal cells, while phosphatidylcholine aids in the repair and replacement of cell membranes.

Anti-inflammatory properties

The anti-inflammatory properties of Phytosome complexes better than their uncomplexed herbal extracts. Phytosome complexes were tested for their anti-inflammatory properties in animals using croton oil-induced dermatitis, an inflammation model that includes both tissue and vascular damage [63]. Results using silymarin, a flavanolignan with anti-phlogistic activity and a free radical scavenger, and glycyrrhetinic acid phytosome from glycyrrhiza glabra, a potent anti-inflammatory and anti-allergic triterpenoid, were reviewed. The ultimate aim was the decrease of edema, and the materials were delivered topically as micro dispersion in water. Although phospholipids by themselves have no anti-inflammatory properties, it was shown that their interaction with glycyrrhetinic acid increased this property's duration. As a result, 18-beta-glycyrrhetinic acid's anti-edematous effect diminished with time and almost stopped after 24 hours, during which its phosphatidyl complex continued to cause an 80% decrease in edema. In the croton oil test in mice, the silymarin complex demonstrated a higher degree of edema reduction (76% after 6 hours) in comparison to that with the free form (33% even after 12 hours) (Fig. 4). Therefore, compared to the free phytoconstituent, the PHYTOSOME was almost twice as effective. The 18-beta-





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glycyrrhetinic acid compound showed strong effectiveness against UVA-induced erythema in a different research[64].

Anti-cancer properties

Alhakamy and a colleague investigated the use of quercetin phytosome functionalized by scorpion venom in the treatment of breast cancer. Pharmacologically active taxifolin (dihydroquercetin) is known for its anti-inflammatory, antibacterial, antiangiogenic, liver-protective, and anticancer characteristics. However, due to its hydrophilic nature, its application was limited. In this work, phospholipids and taxifolin-rich ethyl acetate were mixed, and the resulting Phytosome were tested ex vivo against MCF-7 cell lines. The Phytosome have a lower IC50 value and increased cytotoxic activity[65]. The MCF-7 cell line is susceptible to the anticancer effects of Phytosome loaded with aloe vera extract. Phosphatidylcholine (98% w/w) is the main constituent found in lecithin. Phosphatidylcholine nanoparticles have been shown to have mitogenic effects on MCF-7 breast cancer cells[66]. Specifically, the nanoparticles, when dispersed in a pH 7.0 buffer, alter the plasma membrane of the cancer cells, affecting the proliferation of the cells and the epidermal growth factor receptor (EGFR). Furthermore, Alkamy et al. discovered that Phytosome enriched with thymoquinone have anticancer properties against human lung cancer cells (A549 cell line). The work highlights the significance of this innovative phytosome method supporting cytotoxicity against lung cancer cells and a prolonged release profile[67]. Resveratrol inhibits the action of aromatase, has antiangio genesis and apoptotic properties, and is useful in the treatment of breast cancer. Two complementary anticancer drug combinations from fungal and herbal sources were chosen, synthesized in a multi compartment nano carrier system, and then encapsulated in a resveratrol phytosome for use in this investigation.

Anti aging

The most visible organ, the skin, performs vital tasks including controlling body temperature and sensing pressure, temperature, and pain. It also serves as a vital barrier against harmful substances and pollutants, which makes aging apparent[68-69]. Thinning, sagging, age spots appearing, and dry skin are all signs of aging. Anti-aging solutions are in high demand as a result of the growing desire to look or feel young[70]. Free radicals are extremely reactive oxygen molecules that interact with collagen molecules to cause a loss of tone in the elasticity of the skin, which starts the aging process. Free radical-induced oxidative damage is prevented or slowed down by antioxidants [71]. Antioxidant effects are very helpful in cosmetic compositions. The primary cause of this action in herbs is the redox characteristics of phenolic compounds, which enable them to function as hydrogen donors, reducing agents, and singlet oxygen quenchers[72]. Antioxidant polyphenols found in a wide variety of fruits, vegetables, whole grains, and herbs scavenge and remove free radicals. Natural skincare products are rapidly absorbed by the skin's outer layers and are hypoallergenic in nature[73].

Thus, anti-aging medicines can be produced from plants that contain antioxidants. Orange peel (Citrus auranticum, Rutaceae) extract has excellent skin-whitening, anti-oxidant, and anti-acne properties. Phenolic substances such as flavonoids and phenolic acids are abundant in the peel and seeds of citrus fruits. When compared to seeds, the peels have a higher flavonoid content [74]. Phenolic chemicals, or flavonoids, are present in liquorice (Glycyrrhiza glabra, Fabaceae) [75]. Because phenolic components are present in high concentrations and have considerable ROS scavenging, hydrogen-donating, and reducing properties, liquorice ethanol extract has good anti-oxidant action[76]. It enhances the skin's hydration and viscoelastic qualities[77]. Because the chemical elements in the herbs work synergistically, polyherbal compositions have a high therapeutic impact [78]. Licorice extract and orange peel extract have each been utilized for their possible cosmetic benefits. Thus, an effort was made to investigate how extracts together affect the aging of the skin.

Anti oxidant

In 2006 & 2007, Maiti et al, developed naringenin and curcumin Phytosome in two different studies and reported that antioxidant activity of the phytosomal complex has better therapeutic efficacy. A naturally occurring flavanone with a wide range of biological action is naringenin. Naringenin has to be administered frequently in order to maintain an effective plasma concentration because of its quick clearance. In comparison to free naringenin, we





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assessed the therapeutic potential of the naringenin-phospholipid complex under oxidative stress conditions. At a dosing level of 100 mg kg-1 (p.o.), naringenin-phospholipid complex was produced and its antioxidant activity evaluated in rats intoxicated with carbon tetrachloride. Serum levels of total bilirubin, alkaline phosphatase, glutamate pyruvate transaminase, and glutamate oxaloacetate transaminase were measured in liver function tests. To evaluate the antioxidant capability at the same dose level, indicators for liver function, such as glutathione peroxidase, superoxide dismutase, catalase, and thiobarbituric acid reactive compounds, were evaluated. Naringenin's plasma levels was also evaluated. Compared to free naringenin at the same dosage level, the naringenin-phospholipid complex was found to considerably protect the liver for a longer period of time and to increase the antioxidant activity of the biomolecule. Naringenin's phospholipid complex demonstrated superior antioxidant activity compared to the free compound throughout an extended period of action, potentially decreasing the molecule's rapid excretion from the body[79]. A novel formulation of Curcumin with phospholipid was developed to overcome the limitation of absorption and to investigate the protective effect of curcumin-phospholipid complex on carbon tetrachloride induced acute liver damage in rats.

The antioxidant activity of curcumin-phospholipid complex(Curcumin Phytosome) (equivalent of curcumin 100 and 200 mg/kg body weight) was evaluated by measuring various enzymes in oxidative stress condition. Curcumin-phospholipid complex significantly protected the liver by restoring the enzyme levels of liver glutathione system and that of superoxide dismutase, catalase and thiobarbituric acid reactive substances with respect to carbon tetrachloride treated group (P < 0.05 and < 0.01). The complex provided better protection to rat liver than free curcumin at same doses. Serum concentration of curcumin obtained from the complex (equivalent to 1.0 g/kg of curcumin) was higher (Cmax 1.2 microg/ml) than pure curcumin (1.0 g/kg) (Cmax 0.5 microg/ml) and the complex maintained effective concentration of curcumin for a longer period of time in rat serum. The results demonstrated that, at the same dosage level, curcumin-phospholipid complex exhibits greater hepato protective action than free curcumin due to its superior antioxidant capacity[80]. In two distinct experiments, Maiti et al. synthesized the phytosome of naringenin, a flavonoid from grapefruit, Vitis vinifera, and curcumin, a flavonoid from turmeric, Curcuma longa. In every measured dosage range, the complex's antioxidant efficacy outperformed that of pure curcumin by a significant margin. In the other study, the naringenin phytosome that had been generated outperformed the free compound in terms of antioxidant activity. This difference in antioxidant activity may have been caused by a slower rate at which the molecule was removed from the body[81].

CHARACTERIZATION TECHNIQUES

Particle size determination

The particle size of the phytosome is determined by Dynamic Light Scattering (NANO ZS Malvern instrument) . And Particle size measurement was performed as following 1/100 (v/v) dilution of phytosomal suspension in redistilled water at 25° C[82].

Zeta Potential Determination

Zeta potential is estimated on the basis of the electro phoretic mobility under an electric field. Zeta potential was measured by using the NANO ZS Malvern instrument at 25°C following the same dilution in 1 mM NaCl solution[82].

Percentage yield

The prepared phytosome were dried properly and then weighed accurately. This weight was divided by the total weight of drugs non volatile recipients[82].





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Drug entrapment efficiency

Take 100 mg of prepared Phytosome were dried properly and dispersed it in 50 ml distilled water. The content was stirred for 2 hrs and then filtered through Whatman filter paper of pore size of 45 μ m. Then dilute it and analyzed spectro photometrically at λ max 266. The amount of drug entrapped is calculated from the standard calibration curve[82].

actual amount of drug in

phytosomal formulation
theoretical amount of drug in
phytosomal formulation

Determination of drug content

To determine the drug content of phytosome complex, dissolve accurately weighed 10 mg of a complex in 10 ml methanol. After dilution, the absorbance was determined by UV – Spectrophotometer at 426nm and drug content was determined[83].

Field emission scanning electron microscopy

The Field emission scanning electron microscopy used to determine surface morphology of the Phytosome by using (FE-SEM; FEI Quanta FEG 200, Netherlands). The powdered sample, was placed on a metal stub of an electron magnifying device and coated with gold using a particle spray. A digital representation of the sample was obtained by examining the tip at different magnifications in an irregular manner. Using Image J software, FE-SEM images were used to measure the phytosome diameters randomly[84].

Transmission electron microscopy

The morphological properties of phytosome were assessed through the use of transmission electron microscopy (JOEL Ltd., Tokyo, Japan; TEM-100S Microscope). The samples undergo a 10-minute sonication after being diluted with ethanol at a ratio of 1:20. A carbon-glazed grid was coated with a drop of phytosome, which was then allowed to form a thin film. Using a TEM, phytosome images were captured [84].

Spectroscopic Evaluation of Phytosome

Ultraviolet spectra (UV-spectra)

The Samples that reflect different absorption in the UV wavelength range can be used to characterize own structural properties. Most studies have revealed that there is no differences in the UV absorption characteristics of constituents before and after complexation. Xu et al. prepared the luteolin-phospholipid complexes(Phytosome) and found that the characteristic peaks of luteolin remained present[85]. Therefore, we conclude that the chromophores of the compounds are not affected by being complexed with phospholipids.

Fourier transform infrared spectroscopy (FTIR)

FTIR is a powerful method for the structural analysis, and yields different functional groups that show distinct characteristics in band number, position, shape, and intensity. The formation of the phyto-phospholipid complexes(Phytosome) can be verified by comparing the spectroscopy of phospholipid complexes to that of physical mixtures. The Separate studies may show different results. In fact, Das and Kalita prepared the Rutin Phytosome. The FTIR of a physical mixture of rutin and phyto-phospholipid complexes was superimposable with the pure rutin[86]. When Mazumder et al. prepared the sinigrin-phytosome complexes, the FTIR of phytosome complex showed the different peaks from that of sinigrin, phospholipids and their mechanical mixtures[87].

Differential scanning calorimetry

In DSC, the interactions can be observed by comparing the transition temperature, appearance of new peaks, melting points, disappearance of original peaks, and changes in the relative peaks area. Phyto-phospholipid complexes(Phytosome) usually display the radically different characteristic peaks compared to those of a physical





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mixture. It is concluded that, in addition to the two fatty chains of phospholipids, the strong interactions occur in the active ingredients and the polar part of phospholipids also inhibits free rotation. Das and Kalita prepared the phytophospholipid complexes rutin and the resulting DSC thermogram showed two characteristic peaks that were lower than that of the physical mixture and the peaks of rutin and PC disappeared [88].

X-ray diffraction

At present, the X-ray diffraction is an effective method to examine the microstructure of both crystal materials and some amorphous materials. The X-ray diffraction is usually performed on either active constituents or active constituent phyto-phospholipid complexes, PCs and their physical mixtures. X-ray diffraction of an active constituent and physical mixture shows the intense crystalline peaks that indicate a high crystal form. By contrast, the active constituent phyto-phospholipid complexes do not exhibit the crystalline peak, which suggests that the constituents in complex with phospholipids exhibit a molecular or amorphous form. The conclusion, that phyto-phospholipid complexes exhibit superior hydrophilicity and lipophilicity in comparison to active components[89].

Nuclear magnetic resonance (NMR)

The ¹H NMR and ¹³C NMR techniques plays an very important role in the identification of the structures of the complexes. As mentioned above, interactions between polyphenols and phospholipids are created by the hydrogen bonds rather than chemical bonds. Angelico et al. established based on NMR data that the hydrogen bonds can form between some polar phenolic functional groups of silybin A and phospholipids[90]. The spectra of the different phyto-phospholipid complexes suggest that the hydrophobic side of lipids can act to cover the envelope on the central choline-bioactive parts of these complexes.

Recent Advancements of Phytosome

The virus appeared to spread globally in December 2019 and was declared a pandemic by the World Health Organization (WHO) on March 11, 2020. The SARS-CoV-2 coronavirus is responsible for it, as it has caused an unusually high number of infections and deaths (4.9 million) from acute severe respiratory syndromes across the globe[91]. Quercetin is a flavanol that the human body is not capable of producing. It is slightly soluble in lipids and popular dietary supplement that may offer protection from a number of illnesses. Quercetin phytosome is used as a supplement in COVID-19. The adequate potency of quercetin as a 3CLpro has only recently been confirmed. Recently, quercetin's oral bioavailability in humans has been evaluated by the use of a phospholipid delivery form called Quercetin Phytosome. Pharmacokinetics research revealed that total quercetin had a 20-fold increased bioavailability when its conjugated form, glucuronide, was hydrolyzed. In addition, it was reported that the body can create free systemic quercetin in response to a specific type of glucuronidase. In a therapeutic context, quercetin in a bioavailable form, like Quercetin Phytosome, should be considered as an effective way to treat COVID-19 while also considering its anti-inflammatory and thrombin-inhibitory properties[92]. Mariangela Rondanelli et al

examined the potential impacts of quercetin Phytosome (250 mg twice a day) multivitamin supplements for three months as a defense against symptomatic COVID-19. There are total of 120 individuals (63 men and 57 women, aged 49 and 12) includes there were 60 in the supplements group and 60 in the placebo group. There were no statistically significant differences seen between the groups with regarding gender, smoking, or chronic illness. Participants completed brief COVID-19 diagnostic examinations every three weeks. During the course of our trial, 5 out of 60 subjects in the quercetin group and 4 out of 60 in the control group had COVID-19. Complete clinical recovery was observed at 7 and 15 days in the quercetin and placebo groups, respectively. After five months, the analysis showed that participants who took quercetin supplements had a 99.8% COVID-free survival function (risk of infection), while those in the control group had a 96.5% risk. In order to prevent being infected with COVID-19, patients who took the supplement had a protection factor 14% higher than those who took a placebo, based on the value of EXP(B)[93]. Researchers conduct a lot of research, and the most recent studies show that phytosome technology is an innovative method to improve plant extract absorption and bioavailability while significantly reducing the dose level. Due to





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their possible pharmacological effects, some plant extracts such as silymarin, grape seed extract, quercetin, curcumin, hesperetin, ginkgo biloba extract, Andrographolide, among others are receiving increased attention these days [94-95]. Newer studies have been made possible by the technique's use and the increasing demand for herbal medicines in the current medical environment for the therapy of numerous diseases. Here is a brief summary of some of the important studies done by different researchers. In 1992 & 1994, Schandalik et al, used nine human volunteer patients and tested the hepatoprotective activity of silymarin and reported that the phytosomal form of silybin possess four times greater passage through the liver [96-97]. et al, In 1993, using 232 patients with chronic hepatitis. He also reported the better bioavailability of silymarin phytosome than the uncomplexed form[98]. In 1999, Grange et al, conducted series of experiments on silymarin Phytosome and reported the better fetoprotectant activity of the Silymarin Phytosome. In 2002, Busby et al, also reported the better fetoprotectant activity of silymarin phytosome than uncomplexed silymarin against ethanol-induced behavioural deficit. In 2001, Jhiang et al. Prepared Herba Epimedii flavonoid phytosome (EPF) by using a solvent evaporation approach, and a dissolution study was used to examine the accumulative dissolution of various ratios of EPF-PVP precipitate. According to the study, the precipitate dissolved much more quickly than both the physical mixture and the Herba Epimedii extract tablets [99]. According to Bombardelli et al. (2005), silymarin Phytosome had much greater specific activity and a more enduring effect compared to the individual constituent in terms of the percentage decrease of edema, myeloperoxidase activity inhibition, antioxidant activity, and free radical scavenging activity[100].

Some Patented Technologies Related to Phytosome

There are number of innovative processes and the formulation research studies in the field of Phytosome carried out by the academic scientists as well as in industrial laboratories. Some patents for Phytosome and the other related technologies along with their applications and innovations are listed in Table 1.

CONCLUSION

Phytosome are the novel formulations which offers improved bioavailability of water-soluble herbal constituents through skin and gastrointestinal tract. Phytosome are one of the phospholipid-based drugs delivery system with a better absorption and stability profile as compared to other drug delivery systems. Phytosome have improved pharmacokinetic and pharmacological parameter, which enable them to be used for different therapeutic purposes like cardiovascular, anti-inflammatory, anticancer, immune modulator, antidiabetic etc. Phytosome or herbosomes are advanced and novel form of botanicals and phytoconstituents that are better absorbed both orally, topically and transdermally. The formulation methods and characterization techniques of Phytosome has been well established. Many areas of Phytosome are to be reported in the future in the prospect of pharmaceutical application. The phytosome technology forms a link between the conventional delivery system of phyto constituents and novel drug delivery systems. This review is an attempt to present a concise profile of Phytosome as a delivery system including mechanism of action of phytosome by topical and oral route.

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Table 1: Some patented technologies related to Phytosome.[101]

Title of Patent	Innovation	Patent Number
Phospholipid complexes of olive fruits or leaf extracts having improved bioavailability	Phospholipids complexes of olive fruits or leaf extracts or compositions having improved bioavailability.	EP/1844785
Fatty acid monoesters of sorbityl furfural and compositions for cosmetic and dermatological use	Fatty acid monoesters of sorbityl furfural and compositions for cosmetic and dermatological use.	EP1690862
Treatment of skin and wound repair with thymosin β4.	Compositions and methods for treatment of skin utilizing thymosin $\beta 4$.	US/2007/0015698
An anti-oxidant preparation based on plant extracts for the treatment of circulation and adiposity problems	An anti-oxidant preparation based on plant extracts for the treatment of circulation and adiposity problems Preparation based on plant extracts which has an antioxidant effect and is particularly useful in treatment of circulation problems such as phlebitis, varicose veins, arteriosclerosis, haemorrhoids and high blood pressure.	EP1214084
Compositions comprising gingko biloba derivatives for the treatment of asthmatic and allergic conditions	Compositions containing fractions deriving from ginkgo biloba, useful for the treatment of asthmatic and allergic conditions.	EP1813280
Cosmetic and dermatological composition for the treatment of aging or photo damaged skin	Composition for topical treatment of the skin comprises a substance that stimulates collagen synthesis and a substance that enhances the interaction between extracellular matrix and fibroblasts Cosmetic or dermatological composition for topical treatment.	EP1640041

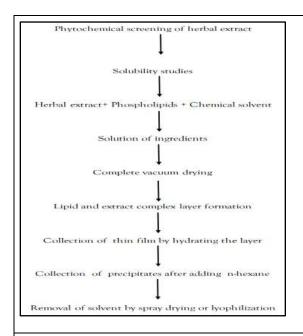


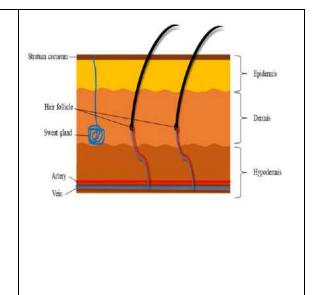


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Soluble isoflavone compositions

Isoflavone compositions exhibiting improved solubility (Example: light transmittance), taste, color, texture characteristics and methods for making.





WO/2004/045541

Figure 1.Common stages in formulation of Phytosome

Transappendageal Routa Intercellular Route Transcellular Route

Figure 2. Simplified structure of the skin and its barriers.

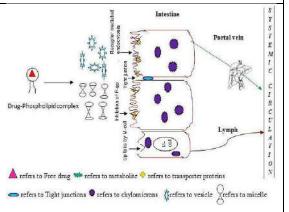


Figure 3. Mechanism of transdermal drug delivery.

Figure 4.Schematic diagram representing the possible mechanism of phyto-phospholipids complex absorption





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RESEARCH ARTICLE

Accurate Equitable Domination in Graphs

P.Nataraj^{1*}, A. Wilson Baskar² and V.Swaminathan³

¹Assistant Professor, Department of Mathematics, Sri Sai Ram Engineering College, (Affiliated to Anna University), Chennai, Tamil Nadu, India.

²Associate Professor, Department of Mathematics, Saraswathi Narayanan College, (Affiliated to Madurai Kamaraj University), Madurai, Tamil Nadu, India.

³Reader (Retd.), Saraswathi Narayanan College, (Affiliated to Madurai Kamaraj University), Madurai, Tamil Nadu, India.

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*Address for Correspondence P.Nataraj

Assistant Professor, Department of Mathematics, Sri Sai Ram Engineering College, (Affiliated to Anna University), Chennai, Tamil Nadu, India. Email: nataraj.maths@sairam.edu.in



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ABSTRACT

A subset $D \subseteq V(G)$ is called an accurate equitable dominating set of G if V - D does not contain an equitable dominating set of cardinality |D|. The minimum cardinality of an accurate equitable dominating set is the called the accurate equitable domination number of G and is denoted by $\gamma_{ae}(G)$. The upper accurate equitable domination number $\Gamma_{ae}(G)$ is the maximum cardinality of a minimal accurate equitable dominating set. In this paper we initiate the study of this parameter.

Keywords: Equitable domination, Accurate domination, Accurate equitable domination.

INTRODUCTION

In a simple graph G a subset S of the vertex set V of G is called a dominating set if a every vertex in the complement of S with respect to V is adjacent with some vertex of S. The cardinality of a minimum dominating set is called the domination number and is denoted by $\gamma(G)$. Two vertices x, y are said to be degree equitable if $|\deg_G(x)|$ $\deg_G(\mathbb{Z}_V) \leq 1$. If $x \in V(G)$, then the open equitable neighbourhood of x consists of those vertices which are adjacent with x and degree equitable with x. The closed equitable neighbourhood of x is the open neighbourhood of xtogether with x. If T is a subset of V then the open equitable neighbourhood of T is the union of the open equitable neighbourhoods of elements of T and the closed equitable neighbourhood of T is the union of the open equitable





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neighbourhood of T and T. For a subset X of V the private equitable neighbour of v with respect to the X consists of those vertices in the closed equitable neighbourhood of v - the closed equitable neighbourhood of $X - \{v\}$. The maximum value of equitable degree of a vertex v in the G is denoted by $\Delta_e(G)$ and the minimum value by $\delta_e(G)$. A subset S of the vertex set V is called an equitable dominating set of G if every vertex in the complement of S with respect to the V is equitably dominated by the some vertex of S. The minimum cardinality of an equitable dominating set of G is denoted by $\gamma_e(G)$. If S is a dominating set of G with the property that the complement of G with respect to the G does not contain any dominating set of cardinality |S|, then G is called an accurate dominating set and the minimum cardinality in referred to as the accurate domination number of G and is denoted by $\gamma_e(G)[7]$.

Definition 1. Suppose *S* is a subset of *V* of *G* such that *S* is an equitable dominating set and V - S does not contain an equitable dominating set of *G* with cardinality |S|. Then *S* is called an accurate equitable dominating set of *G* and its minimum cardinality is called the accurate equitable domination number of *G* and is denoted by $\gamma_{ae}(G)$. The maximum cardinality of a minimal accurate equitable dominating set of *G* is denotes by $\Gamma_{ae}(G)$.

$\gamma_{ae}(G)$ Of Some Standard Graphs

i)
$$\gamma_{ae}(K_n) = \left|\frac{n}{2}\right| + 1$$
, for $n \ge 3$
ii) $\gamma_{ae}(P_n) = \begin{cases} \frac{n}{3}, & n \equiv 0 \pmod{3} \\ \left[\frac{n}{3}\right] + 1, & n \equiv 1 \text{ or } 2 \pmod{3} \end{cases}$
iii) $\gamma_{ae}(C_n) = \begin{cases} \left[\frac{n}{3}\right] + 2, & n \equiv 0 \pmod{3}, \\ \left[\frac{n}{3}\right] + 1, & n \ne 0 \pmod{3} \end{cases}$
iv) $\gamma_{ae}(W_n) = \begin{cases} 3, & \text{if } n = 4 \\ 1, & \text{if } n = 5 \\ \gamma_{ae}(C_{n-1}) + 1, & \text{if } n \ge 6 \end{cases}$
v) $\gamma_{ae}(K_{m,n}) \begin{cases} min(m,n), if |m-n| \le 1 \\ m+n, if |m-n| \ge 2 \end{cases}$
vi) $\gamma_{ae}(K_{1,n}) = n+1$, for $n \ge 3$

Bounds On $\gamma_{ae}(G)$

Remark 1. Every accurate equitable dominating set is an equitable dominating set.

Remark 2. An accurate equitable dominating set may or may not be a minimal equitable dominating set.

Proposition 1. For any graph G, $\gamma_e(G) \leq \gamma_{ae}(G)$.

Proposition 2. If G contains an equitable isolate, then a minimal equitable dominating set of G is an accurate equitable dominating set of G.

Proof. Let v be an equitable isolate of G. Then, $v \in D$ for every equitable dominating set D of G. Thus, |V - D| cannot have an equitable dominating set of cardinality D.

Corollary 1. If *G* contains an equitable isolate, then $\gamma_{ae}(G) = \gamma_e(G)$

Proposition 3. For any graph G, $\gamma_{ae}(G) \leq n - \gamma_e(G) + 1$

Proof. Let *D* be a minimum equitable dominating set of *G*. Then, for any $v \in V - D$, $(V - D) \cup \{v\}$ is an accurate equitable dominating set of *G*. Hence, $\gamma_{ae}(G) \le |(V - D) \cup \{v\}| \le n - \gamma_e(G) + 1$.





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Corollary 2. If *G* contains equitable isolates, then $\gamma_{ae}(G) \leq \frac{n+1}{2}$

Proposition4. Let G be a simple graph with maximum equitable degree Δ_e , then $\frac{n}{\Delta_e+1} \leq \gamma_{ae}(G) \leq \frac{n\Delta_e}{\Delta_e+1}+1$

Proof. Let D be a γ_e - set of G. Each vertex $v \in V(G)$ can equitable dominate at most $\Delta_e + 1$ vertices. Hence, $\gamma_e(G) \ge \frac{n}{\Delta_e + 1}$. Now, since $\gamma_e(G) \le \gamma_{ae}(G)$, the lower bound holds. Now by **Proposition 3**

$$\begin{array}{l} \frac{\Delta_e + 1}{\gamma_{ae}(G)} \le n - \gamma_e(G) + 1 \\ \le n - \frac{n}{\Delta_e + 1} + 1 \end{array}$$

$$\leq \frac{n\Delta_e}{\Delta_e+1}+1$$

Proposition 5. For any graph G with $n \ge 2$, an equitable dominating set D such that $|D| = \left\lfloor \frac{n}{2} \right\rfloor + 1$, then D is an accurate equitable dominating set.

Proof. Let D be an equitable dominating set with $|D| = \left\lfloor \frac{n}{2} \right\rfloor + 1$. Then $|V - D| \le \frac{n}{2}$. Hence D is an accurate equitable dominating set.

Proposition 6. Let *T* be a tree with *s* cut vertices. Then, $\gamma_{ae}(T) \le s + 1$. Equality holds if and only if each cut vertex is adjacent to a pendent vertex.

Proof. Let S be the set of all cut vertices of T with |S| = s. Let u be a pendent vertex of G. Then $S \cup \{u\}$ is an accurate equitable dominating set of T.

Suppose $\gamma_{ae}(T) = s + 1$. If there exists a cut vertex $v \in S$ such that v is not adjacent to any of the end vertices, then S is an accurate equitable dominating set, a contradiction. Converse is obvious.

Definition 2 [9] For a simple graph G with the vertex set V(G), a subset S of V(G) is called a maximal equitable dominating set of G, if the complement of S with respect to the V not an equitable dominating set of G. If such a set has a minimum cardinality then that cardinality is called the maximal equitable domination number of G and is denoting $Y_e^m(G)$.

Proposition 7. For any graph G, $\gamma_{ae}(G) \leq \gamma_e^m(G)$, equality holds if $G = C_n$.

Definition 3 [1] Suppose *S* is a subset of the vertex set *V* of a simple graph *G*, *S* is called an equitable total dominating set of *G* if *S* is an equitable dominating set of *G* and the subgraph included by *S* has no isolates. The minimum cardinality of such a set is called the equitable total domination number of *G* and is denoted by $\gamma_e^t(G)$.

Proposition 8. If a simple graph *G* has a pendent vertex, then $\gamma_{ae}(G) \leq \gamma_e^t(G) + 1$.

Proof. Let G be a graph and let $v \in V(G)$ be a pendent vertex and $u \in V(G)$ be its support vertex. Let D be a γ_e^t - set of G.

Case 1: Suppose $v \in D$. Then, u is also in D. Hence D is an accurate equitable dominating set of G. Thus, $\gamma_{ae}(G) \le |D| = \gamma_e^t$.

Case 2: Suppose $v \notin D$. Then, $u \in D$. Hence $D \cup \{v\}$ is an accurate equitable dominating set of G. Thus, $\gamma_{ae}(G) \leq |D \cup \{v\}| = \gamma_e^t + 1$.





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When $G = P_4$, $\gamma_{ae}(P_4) = 3 = 2 + 1 = \gamma_e^t + 1$.

Graphs With $\gamma_{ae} = \gamma_e$

Definition 4. Suppose H is a simple graph. Suppose one pendent vertex is attached with every vertex of H then the resulting graph is called the corona of H and is denoted by $H \circ K_1$.

Proposition 9. Let G = (V, E) be a graph. Then $\gamma_{ae}(G) = \gamma_e(G)$ if and only if there exists a set $S \in D_{\gamma_e}(G)$ such that $S \cap S' \neq \emptyset$ for every set $S' \in D_{\gamma_e}(G)$, where $D_{\gamma_e}(G) = \{S \mid S \text{ is a } \gamma_e - \text{set } \}$.

Proof. Let G be a graph with $\gamma_{ae}(G) = \gamma_e(G)$. Let S be a minimum accurate equitable dominating set of G. Since S is an equitable dominating set and $|S| = \gamma_{ae}(G) = \gamma_e(G)$, $S \in D_{\gamma_e}$. Let S' be an arbitrary minimum equitable dominating set of G. If $S \cap S' = \emptyset$, then $S' \subseteq V(G) - S$, which implies S' is an equitable dominating set of G of cardinality |S|, a contradiction. Hence, $S \cap S' \neq \emptyset$.

Conversely, let $S \in D_{\gamma_e}(G)$ such that $S \cap S' \neq \emptyset$ for every set $S' \in D_{\gamma_e}(G)$. Then, S is an accurate equitable dominating set of G. Therefore, $\gamma_{ae}(G) \leq |S| = \gamma_e(G) \leq \gamma_{ae}(G)$. Thus, we must have $\gamma_{ae}(G) = \gamma_e(G)$.

Corollary 3. If $G = H \circ K_1$, $H \neq K_1$, then $\gamma_{ae}(G) = \gamma_e(G)$

CONCLUSION

In this paper we initiated the study of a new parameter accurate equitable domination number of graph and obtained some results. More results on this parameter can be obtained using other domination parameters.

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RESEARCH ARTICLE

Anti-Hyperglycemic Effect of Siddha Varmam Formulation Neerizhivu Kudineer in Streptozotocin Induced Diabetics Rats

Thirunarayanan G^{1*}, Suvedha P² and Jayalakshmi J³

¹Medical Officer, National Institute of Siddha, Tambaram Sanatorium, Chennai, Tamil Nadu, India.

²PG Scholar, Department of Kuzhandhai Maruthuvam, National Institute of Siddha, Tambaram Sanatorium, Chennai, Tamil Nadu, India.

³Professor, Sivaraj Siddha Medical College, Salem and Ph.D Scholar (Part Time), National Institute of Siddha, Chennai, Tamil Nadu, India.

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*Address for Correspondence G.Thirunarayanan,

Medical Officer, National Institute of Siddha, Tambaram Sanatorium, Chennai, Tamil Nadu, India.

Email: thirunarayanan.dr@gmail.com



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ABSTRACT

Varmam is a part of the Siddha system of medicine and it is well known for its therapeutic action in various health conditions. Varmam can work well in various lifestyle disorders like diabetes and hypertension. These life style disorders can be treated with both Varmam internal medicine and Varmam therapies. The Trail drug Neerizhivu Kudineer is an herbal formulation quoted in ancient Varma text for Neerizhivu (Diabetes mellitus). The study aims to determine the anti-hyperglycemic effect of Siddha Varmam formulation Neerizhivu Kudineer in streptozotocin induced diabetic rats. Neerizhivu Kudineer were analysed for anti-Hyperglycemic activity at the doses of 100 and 200 mg/kg orally continue for 28 days, with reference to the fasting blood glucose (FBG) and blood glycosylated haemoglobin (HbA1c) in the STZ induced diabetic. other factors like triglyceride (TG), total cholesterol (TC), serum biochemical parameters (AST, ALT, ALP), lipid peroxidation (MDA), HDL, total protein, body weight and liver antioxidants enzymes (GSH, SOD, CAT) levels were also analyzed in the STZ induced diabetic rats. The results of this study reveal that the traditional Varmam internal medicine Neerizhivu Kudineer has an antihyperglycemic effect and it can help in the reduction of blood glucose levels in patients suffering from diabetes mellitus.

Keywords: Neerizhivu Kudineer, Neerizhivu noi, Diabetes mellitus, Varma Maruthuvam, Siddha system of medicine.





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INTRODUCTION

Varmam Maruthuvam is part Siddha system of medicine and it is one of the oldest traditional medicines in the world. Varmam Medicines are widely used in southern parts of India, predominantly in Tamilnadu[1]. Neerizhivu is a disease that has been given much importance both in the past and the present. Neerizhivu is a clinical or health condition characterized by frequent urination with more volume, invasion of ants and house flies at the place of urination, smelly urine with a sweet odour while it gets boiled, gradual loss of weight with the reduction in power of the human body and seven thathus[2]. The symptoms of Neerizhivu can be correlated with Diabetes in modern terms. Certain symptoms such as a burning sensation in the palm and soles, delayed wound healing etc., are the classic indicators of this Neerizhivu (Diabetes mellitus). Diabetes currently affects more than 62 million Indians, which is more than 7.1% of the adult population [3]. The average age of onset of Neerizhivu Noi is 42.5 years. Nearly one million Indians die due to Neerizhivu Noi every year[4]. According to the Indian Heart Association, India is projected to be home to 109 million individuals with Neerizhivu No by 2035[5]. Siddha and Varmam Maruthuvam have various herbal and herbo mineral formulations which are giving remarkable changes in the management of Neerizhivu (Diabetes mellitus). One among these herbal formulations is Neerizhivu Kudineer mentioned in Varmam literature. The ingredients in Neerizhivu Kudineer have been used in the treatment of diabetes mellitus by common people as home remedies. Even though many herbal formulations are being used for its therapeutic action in day-to-day practices, it also has to be evaluated scientifically. Hence here it is tried to evaluate the anti-hyperglycemic effect of Siddha Varmam formulation Neerizhivu Kudineer in streptozotocin-induced diabetics rats.

MATERIALS AND METHODS

Ingredients of *Neerizhivu Kudineer* [6, 7] Show the table No. 1: Drug Description

Collection, Identification and Authentication of the Drug

The drugs mentioned in the Table No.1 were collected from Tirunelveli and Kanyakumari district, Tamilnadu, India. Collected raw drugs were identified and authenticated by Botanist of Government Siddha Medical College, Palayamkottai, Tirunelveli, Tamilnadu.

Purification and Preparation of the Drug

The ingredients of this *Neerizhivu Kudineer* were purified according to the proper methods described in Siddha Classical Literature.

Animals

Healthy Wistar albino rats (weighing 185±10 g) of either sex were used in the study. Rats were kept in standard laboratory conditions i.e., 12 h light/dark cycles at 25-28 °C, relative humidity 55–60 % and were fed with a standard pellet diet and water. The animals were acclimatized to laboratory conditions for one week before the experiment.

Drugs and Chemicals

Streptozotocin was purchased from HI Media Laboratories Pvt. Ltd. India and glibenclamide (reference drug) was from Sanofi India Ltd. India. Trichloroacetic acid (TCA) from Merck Ltd., Mumbai, India; thiobarbituric acid (TBA), 5,5′-dithio bis-2-nitro benzoic acid (DTNB), phenazonium methosulfate (PMS), nicotinamide adenine dinucleotide (NADH) and reduced glutathione (GSH) from SISCO Research Laboratory, Mumbai, India; potassium dichromate, glacial acetic acid from Ranbaxy, Mumbai; and all the other reagents kits used were from Span Diagnostics Ltd. India.





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Oral Glucose Tolerance Test (OGTT)

The OGTT was performed in overnight fasted Wistar albino rats. Rats were divided into three groups (n=6). Group I served as normal control (received distilled water 5 ml/kg b.w. p.o.), groups II and III received Siddha preparation *Neerizhivu Kudineer* at the doses of 100 and 200 mg/kg b.w. p.o. respectively. All the animals received glucose (2 g/kg b.w., orally) 30 min after drugs administration. Blood sample was withdrawn from the tail vein at 0, 30, 60, 120 and 240 min. after glucose administration[8]and blood glucose level was measured using a single touch glucometer (Contour TS, Bayer Health Care.).

Induction of Diabetes in Rats

The rats were rendered diabetic by a single intraperitoneal dose of 50 mg/kg b.w. STZ freshly dissolved in ice cold 0.1 M citrate buffer (pH 4.5)[9]. After 72 h, fasting blood glucose (FBG) levels were measured and only those animals showing blood glucose levels≥ 250 mg/dl were used for the present investigation [10]. The day on which hyperglycemia had been confirmed was designated as day 0 [11].

Treatment Schedule and Estimation of Fasting Blood Glucose (FBG) Level

After induction of experimental diabetes, the rats were divided into five groups (n=6). Except for group I, which served as normal (non-diabetic) control, all other groups were comprised of diabetic rats. Group II served as diabetic (STZ) control. Groups III and IV received *Neerizhivu Kudineer* (100 and 200 ml/kg b.w., p.o. respectively) and group V received reference drug glibenclamide (0.5 mg/kg b.w., p.o) daily for 28days [12].

Treatment Protocol

Group I (Normal control): Normal Rats + Normal saline (5 ml/kg b.w.)

Group II (Diabetic control): Diabetic Rats + Normal saline (5 ml/kg b.w.)

Group III (Test low dose): Diabetic Rats + Neerizhivu Kudineer (100 ml/kg b.w.)

Group IV (Test high dose): Diabetic Rats + Neerizhivu Kudineer (200 ml/kg b.w.)

Group V (Standard): Diabetic Rats + Glibenclamide (0.5 mg/kg b.w.)

FBG level was measured on days 0, 14, 21 and 28by using a one-touch glucometer. After 24 hours of the last dose and 18 hours of fasting, blood was collected from all rats in each group by cardiac puncture for estimation of glycosylated hemoglobin (HbA1C), serum lipid profile and serum biochemical parameters and then the animals were sacrificed for collection of liver tissue to check the different endogenous antioxidant parameters.

Body Weight

The body weight of rats from each group was measured on days 0, 14, 21 and 28. Weight was measured using standard digital weight balance to get accuracy.

Estimation of Glycosylated Hemoglobin (HbA1c)

Glycosylated hemoglobin was analyzed by using a commercially available kit (Beacon Diagnostic Pvt. Ltd. India) which is based on the principle of the ion exchange resin method.

Estimation of Liver Biochemical Parameters and Antioxidant Status

All animals were sacrificed after 24 hr of the last dose by overdose of diethyl ether. Blood samples were collected and the liver was excised, washed in ice-cold phosphate buffer saline, blotted and weighed. 10% w/v of liver homogenate was prepared in 0.15 M Tris-HCl buffer (pH 7.4). The homogenate was centrifuged at 2000×g for 20 minutes at 4°C to remove the cell debris and then the supernatant was centrifuged (REMI C-24) at 12000×g for 1 hour at 4°C. The supernatant was obtained and used for the determination of Lipid peroxidation and endogenous antioxidant enzymes like superoxide dismutase (SOD), catalase (CAT) and non-enzymatic antioxidant i.e. reduced glutathione (GSH).





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Estimation of Lipid Peroxidation Level (TBARS)

Lipid peroxidation (LPO) was assayed according to the method of Ohkawa et al. (1979). To 1 ml of tissue homogenate, 1 ml of normal saline (0.9%, w/v) and 2.0 ml of 10% trichloroacetic acid were added and mixed well. The mixture was then centrifuged (3000 × g) at room temperature for 10 min to separate proteins. 0.5 ml 1.0%thiobarbituric acid was added to 2 ml of supernatant followed by heating at 95 °C for 60 min to generate the pink-colored malondialdehyde (MDA). The absorbance of the samples was measured at 532 nm using a spectrophotometer (Beckman DU 64). The level of lipid peroxides i.e. thiobarbituric acid reactive substances (TBARS) was estimated and expressed as mM of MDA/mg wet tissue using the extinction coefficient of 1.56 ×105M–1 cm -1.

Superoxide Dismutase (SOD) Activity Assay

Superoxide dismutase (SOD) activity was assayed according to the method of Murklund and Murklund[14]. The liver homogenate was prepared in Tris buffer and centrifuged for 40 min at 10,000 r.p.m. at 4 °C. The supernatant was used for the enzyme assay. 2.8 ml Tris–EDTA (composed of 49.78 mM Tris, 0.0012 mM EDTA; pH 8.5) and 100 μ l pyrogallol (2 mM) were taken in the cuvette and measured the auto-oxidation of pyrogallol by measuring the absorbance for 3 min (reading taken at 30 s interval for 3 min) at 420 nm wavelength. Then 2.8 ml Tris– EDTA (pH –8.5), 100 μ l pyrogallol and 50 μ l tissue homogenate were taken and scanned for 3 min at the same wavelength. One unit of SOD activity is the amount of the enzyme that inhibits the rate of auto-oxidation of pyrogallol by 50% and was expressed as Units/mg protein/min. The enzyme unit can be calculated by using the following equations: [50% inhibition =1 U].

Estimation of Catalase (CAT) Activity

Catalase activity was measured based on the ability of the enzyme to break down H2O2. Catalase (CAT) was assayed and expressed as moles of H2O2 decomposed KU/min/mg of tiss[13]. A 10 μ l sample was taken in a tube containing 3 ml of H2O2 in phosphate buffer (M/15 phosphate buffer; pH –7.0). The time required for 0.05 optical density changes was observed at 240 nm against a blank containing the enzyme source in H2O2-free phosphate buffer. One unit of CAT activity is the amount of enzyme that liberates half the peroxide oxygen from H2O2 solution of any concentration in 100 s at 25 °C which is determined by CAT activity expression:

KU of H2O2 consumed/min (units/mg)

E is the optical density at 240 nm and is the time required for a decrease in the absorbance.

Estimation of Reduced Glutathione (GSH) Level

Non-enzymatic antioxidant i.e. reduced glutathione (GSH) was determined by the method of Ellman (1959)[15]. GSH was estimated spectrophotometrically by the determination of DTNB (dithiobis-2-nitrobenzoic acid) reduced by SH groups, expressed as μ g/mg wet tissue. To 0.1 ml of tissue homogenate, 2.4 ml of 0.02 M EDTA solution was added and kept in an ice bath for 10 min. Then 2 ml of distilled water and 0.5 ml of 50% (w/v) trichloro acetic acid were added. This mixture was kept on ice for 10–15 min and then centrifuged at 3000 ×g for 15 min. To 1 ml of supernatant, 2.0 ml of Tris buffer (0.4 M) was added. Then 0.05 ml of DTNB solution (Ellman's reagent; 0.01 M DTNB in methanol) was added and vortexed thoroughly. OD was read (within 2–3 min after the addition of DTNB) at 412 nm in a spectrophotometer against a reagent blank. Appropriate standards were run simultaneously.

Estimation of Serum Biochemical Parameters

Collected blood was analyzed for various Serum biochemical parameters like Serum Aspartate transaminase (AST), Alanine transaminase (ALT) and Serum alkaline phosphatase (ALP). Total protein and lipid profiles. All the analyses were performed by using commercially available kits from Span Diagnostics Ltd. India.

Statistical Analysis

All the results are shown as mean ±SEM. The results were analyzed for statistical significance by one-way analysis of variance (ANOVA) followed by NewmannKeuls multiple range tests. using Graph Pad Prism 5.0 software (Graph Pad Software, USA). P values of < 0.05 were considered as statistically significant.





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RESULTS

Oral Glucose Tolerance Test

Table 2 shows glucose loading to the normal rats increased blood glucose levels from 68 ± 1.8 to 169 ± 3.45 at 60 min and returned to normal at 240 min. Administration of *Neerizhivu Kudineer* at the doses of 100 and 200 ml/kg significantly (p < 0.01) improved glucose tolerance in a dose-dependent manner at 120 min.

Body Weight

The body weights of normal and diabetic rats were summarized in Table 3. The final body weights were significantly (p < 0.01) decreased in the diabetic control group as compared to a normal control group. Administration of *Neerizhivu Kudineer* at the doses of 100 and 200 ml/kg significantly (p < 0.01) improved the body weight when compared to the diabetic control group.

Fasting Blood Glucose Level

There was a significantly (p < 0.01) elevated FBG level in STZ-induced diabetic rats as compared to the normal control group. Administration of *Neerizhivu Kudineer* in diabetic rats at the doses of 100 and 200 ml/kg significant (p < 0.01) reduced the FBG level towards normal as compared to the diabetic control group (Table 4).

Estimation of Serum Lipid Profiles

Serum lipid profiles like total cholesterol and triglyceride in STZ-induced diabetic rats were significantly (p < 0.001) elevated and the HDL level significantly (p < 0.01) decreased compared to the normal control group. Treatment with Neerizhivu Kudineer at the doses of 100 and 200 ml/kg significantly (p < 0.01) reduced the total cholesterol and triglyceride levels and significantly (p < 0.01) increased the HDL level when compared to the diabetic control group (Table 5).

Estimation of Glycosylated Hemoglobin (HbA1c)

Glycosylated haemoglobin level in STZ-induced diabetic rats was significantly (p < 0.01) increased compared to the normal control group. Treatment with *Neerizhivu Kudineer* at the doses of 100 and 200 ml/kg significantly (p < 0.01) reduced the HbA1c level when compared to the diabetic control group (Table 6).

Estimation of Serum Biochemical Parameters

Biochemical parameters like AST, ALT and SALP in STZ-induced diabetic rats were significantly (p < 0.01) elevated and the total protein content was significantly (p < 0.001) decreased compared to the normal control group. Treatment with *Neerizhivu Kudineer* at the doses of 100 and 200 ml/kg significantly (p < 0.001) reduced the AST, ALT and SALP levels and significantly (p < 0.01) increased the total protein level as compared to the diabetic control group (Table 7)

Estimation of Tissue Antioxidant Parameter

Lipid peroxidation results in the formation of ROS species and subsequently elevates the level of malondialdehyde (MDA) in the liver tissue of STZ-induced diabetic rats. In the present study the MDA level was significantly (p < 0.01) increased in STZ-induced diabetic rats compared to the normal control group. Interestingly, treatment with *Neerizhivu Kudineer* at the doses of 100 and 200 ml/kg significantly (p < 0.01) reduced the MDA levels compared to the diabetic control group (Table 8). The levels of catalase reduced GSH and SOD were significantly (p < 0.001) decreased in STZ-induced diabetic rats compared to the normal control group. Administration of *Neerizhivu Kudineer* at the doses of 100 and 200 ml/kg significantly (p < 0.01) increased GSH, SOD and catalase levels in the liver of STZ-induced diabetic rats compared to the diabetic control group (Table 8).





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DISCUSSION

This study is aimed to investigate the anti-hyperglycemic activity of *Varmam* preparation *Neerizhivu Kudineer* in STZ-induced diabetic rats. It was observed that there was significant increase in the fasting blood glucose (FBG), blood glycosylated haemoglobin (HbA1c), triglyceride (TG), total cholesterol (TC), serum biochemical parameters (AST, ALT, ALP), lipid peroxidation (MDA) and significantly decreased in the HDL, total protein, body weight and liver antioxidants enzymes (GSH, SOD, CAT) levels in the STZ induced diabetic rats when compared to normal control group. The *Varmam* preparation *Neerizhivu Kudineer* has shown the presence of alkaloids, flavonoids, steroids and triterpenoids when phytochemical screening was done. Hence, the present study was carried out to evaluate the antidiabetic activity of the *Varmam* preparation *Neerizhivu Kudineer* on streptozotocin-induced diabetic rats. Here *Varmam* preparation of *Neerizhivu Kudineer* at doses of 100 and 200 ml/kg orally continued for 28 days on streptozotocin-induced diabetic rats has revealed that the increase in blood glucose level was observed normal in glucose-loaded rats than in *Neerizhivu Kudineer* treated rats during oral glucose tolerance test (OGTT) and significantly reduced the glycosylated haemoglobin(HbA1c). Diabetic rats treated with *Neerizhivu Kudineer* showed significant improvement in body weight, further there was a significant reduction in serum triglycerides, total cholesterol and an increase in HDL.

Twenty-Eight days of treatment with *Varmam* preparation *Neerizhivu Kudineer* has restored the serum hepatic biochemical parameters toward the normal values in a dose-dependent manner, thereby alleviating liver damage caused by STZ-induced diabetes and also inhibited hepatic lipid peroxidation revealed by the reduction of TBARS levels towards normal with significant increase in the hepatic SOD, catalase and GSH activities in the diabetic rats. This result clearly shows that *Varmam* preparation *Neerizhivu Kudineer* contains a free radical scavenging activity, which could exert a beneficial action against pathological alteration caused by the presence of superoxide radicals and hydrogen peroxide radicals. It is observed that *Neerizhivu Kudineer* treated group has normalized elevated blood glucose level, glycosylated haemoglobin and body weight, further, it restored serum and liver biochemical parameters towards normal values when compared to the diabetic control group. These activities of *Varmam* preparation *Neerizhivu Kudineer* are due to the presence of the above-mentioned phyto constituents and it is assumed that the *Varmam* preparation *Neerizhivu Kudineer* may be responsible for the stimulation of insulin release and also for the observed restoration of metabolic activity.

CONCLUSION

This study reveals the anti-hyperglycaemic action of *Varmam* medicine *Neerizhivu Kudineer*. *Neerizhivu Kudineer* has a favourable effect not only on blood glucose levels but also on glycosylated haemoglobin, serum lipids, antioxidants, serum biochemical parameters and body weight. This *Varmam* preparation *Neerizhivu Kudineer* can act as an antidiabetic agent in treating diabetic complications.

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Table No. 1: Drug Description

S.No	Drugs	Botanical Name	Family	Parts Used
1.	AlamVithai	Ficus benghalensis	Moraceae	Seed
2.	VilamPisin	Limonia acidissima	Rutaceae	Gums

Table 2.Effect of Neerizhivu Kudineer on Blood glucose(mg/dl) in oral glucose tolerance test (OGTT).

Groups	0 min	30 min	60 min	120 min	240 min
Normal control	68 ± 1.8	112 ± 2.25	169± 3.45	126 ± 2.36	72.75 ± 1.95
Neerizhivu Kudineer 100mg/kg	61 ± 1.75	90 ± 1.98a**	95 ± 2.05 a**	88.45 ± 2.46 a**	65.47 ± 2.33 a**
Neerizhivu Kudineer 200mg/kg	52.25 ± 1.38	84.15 ± 2.12a**	69.28 ± 1.35 a**	62.20± 1.24 a**	58.10± 1.02 a**

Each value is expressed as Mean \pm SEM, where n = 6. a Normal control vs. all treated group, **P < 0.01

Table 3. Effect of Neerizhivu Kudineer on fasting blood glucos(mg/dl) level in STZ-induced diabetic rats.

Groups	Before STZ	Day 0	Day 14	Day 21	Day 28
Normal control	75.34 ± 2.14	76.90 ± 2.45	70.70 ± 1.85	65.75 ± 1.56	63.65 ±1.30
Diabetic control (STZ 50 mg/kg)	72.35 ± 1.97	345.20 ± 6.38a**	360.10 ± 6.55 a**	376 ± 6.69 a**	385.25 ± 6.85 a**
Diabetic + Neerizhivu Kudineer (100 ml/kg)	68.74 ± 1.88	324.30 ± 6.18b**	224.45 ± 5.86 b**	116.40 ± 2.65 b**	98.36 ± 2.06 b**
Diabetic + Neerizhivu Kudineer (200 ml/kg)	76.28 ± 1.62	314.46 ± 5.75 b**	175.45 ± 3.60 b**	103.28 ± 1.95 b**	84.55 ± 1.22 b**
Diabetic + Glibenclamide (0.5 mg/kg)	73.37 ± 1.08	290.15 ± 5.45 b**	150.30 ± 1.65 b**	95.75 ± 2.08 b**	66.54 ± 1.63 b**

Each value is expressed Mean \pm SEM, where n=6. a Normal control group vs. diabetic control group, b Diabetic control group vs. all treated group, **p < 0.001.

Table 4. Effect of Neerizhivu Kudineer on body weight (grams) in STZ- induced diabetic rats.

Groups	Day 0	Day 14	Day 21	Day 28
Normal control	175 ± 3.28	172.5 ± 4.80	175.6 ± 5.12	180.4 ± 5.40
Diabetic control (STZ 50 mg/kg)	160 ±3.15	158.5 ± 4.20 a**	142.5 ± 3.80 a**	140.4 ± 3.70 a**





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Diabetic + Neerizhivu Kudineer	162.5 ± 3.22	158.20± 3.34 b**	155.10 ± 3.45 b**	152.20±3.26 b**
(100 ml/kg)				
Diabetic + Neerizhivu Kudineer	164.2 ± 3.75	161.30± 3.48 b**	158.7 ± 2.92 b**	156.6 ± 2.78 b**
(200 ml/kg)				
Diabetic + Glibenclamide	170.2 ± 3.45	166.25± 3.36 b**	163.20± 3.45 b**	161.2 ±3.28 b**
(0.5 mg/kg)				

Each value is expressed as Mean \pm SEM, where n=6. a Normal control group vs. diabetic control group, b Diabetic control group vs. all treated group, **p < 0.001.

Table 5. Effect of Neerizhivu Kudineer on serum lipid profiles (mg/dl)in STZ-induced diabetic rats.

Groups	Total cholesterol	Triglyceride	HDL
Normal control	45.90 ± 3.20	66.75 ± 3.60	64.30 ± 3.40
Diabetic control (STZ 50 mg/kg)	112.7 ± 4.63 a**	155.1 ± 4.95 a**	30.15 ± 1.75 a**
Diabetic + Neerizhivu Kudineer(100 ml/kg)	53.45 ± 2.53 b**	86.85 ± 2.86b**	43.35 ± 1.78 b**
Diabetic + Neerizhivu Kudineer(200 ml/kg)	49.80 ± 2.25b**	74.35 ± 2.67b**	55.30 ± 1.85b**
Diabetic + Glibenclamide	47.25 ± 1.80 b**	70.30 ± 2.55 b**	60.35 ± 1.96 b**
(0.5 mg/kg)			

Each value is expressed as Mean \pm SEM, where n=6. a Normal control group vs. diabetic control group, b Diabetic control group vs. all treated group, **p < 0.001.

Table 6. Effect of Neerizhivu Kudineer on glycosylated hemoglobin in STZ-induced diabetic rats.

Groups	HbA1c (%)
Normal control	7.38 ± 0.85
Diabetic control (STZ 50 mg/kg)	16.40 ±1.70 a**
Diabetic + Neerizhivu Kudineer(100 ml/kg)	9.15 ± 1.45b**
Diabetic + Neerizhivu Kudineer(200 ml/kg)	8.24 ± 1.36 b**
Diabetic + Glibenclamide(0.5 mg/kg)	$7.70 \pm 1.27 b^{**}$

Each value is expressed as Mean \pm SEM, where n=6. a Normal control group vs. diabetic control group, b Diabetic control group vs. all treated group, **p < 0.001.

Table 7. Effect of Neerizhivu Kudineer on serum biochemical parameters and total protein in STZ induced diabetic rats.

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Groups	AST (IU/L)	ALT (IU/L)	ALP (IU/L	Total protein (g/dl)
Normal control	86.20 ± 2.75	73.95 ± 2.60	4.25 ± 0.55	8.14 ± 0.38
Diabetic control (STZ 50 mg/kg)	126.95 ± 3.45 a**	128.15 ± 3.65 a**	16.32 ± 1.05 a**	$4.28 \pm 0.52 \ a^{**}$
Diabetic + Neerizhivu Kudineer	108.66 ± 3.12b**	101.65 ± 2.55 b**	7.70 ± 0.69 b**	6.80± 0.58 b**
(100 ml/kg)				
Diabetic + Neerizhivu Kudineer	96.90 ± 2.36b**	88.85 ± 2.24 b**	$5.05 \pm 0.60 \text{ b**}$	$7.12 \pm 0.68 b^{**}$
(200 ml/kg)				
Diabetic + Glibenclamide(0.5 mg/kg)	92.65 ± 2.16 b**	82.60± 1.85 b**	4.48 ± 0.55 b**	7.56± 0.76 b**

Each value is expressed as Mean \pm SEM, where n=6. a Normal control group vs. diabetic control group, b Diabetic control group vs. all treated group, **p < 0.001.





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Table 8. Effect of *Neerizhivu Kudineer* on lipid peroxidation (MDA), reduced glutathione (GSH) superoxide dismutase (SOD)and catalase (CAT) in STZ- induced diabetic rats.

Groups	MDA	GSH	SOD	CAT
	(mM of MDA/mg	(µg GSH/mg	(mU/min/mg tissue)	(KU/min/mg
	tissue)	tissue)		tissue)
Normal control	92.15 ± 2.60	3.55 ± 0.26	2.78 ± 0.13	4.15 ± 0.36
Diabetic control (STZ 50 mg/kg)	158.90 ± 4.12 a**	1.75 ± 0.14 a**	$1.45 \pm 0.12 \ a^{**}$	$1.80 \pm 0.15 \ a^{**}$
Diabetic + Neerizhivu Kudineer	128.40 ± 3.15b**	2.72 ± 0.22 b**	1.90 ± 0.09 b**	2.60 ± 0.26 b**
(100 ml/kg)				
Diabetic + Neerizhivu Kudineer	115.90 ± 3.45 b**	2.95 ± 0.18 b**	$2.40 \pm 0.16 \ b^{**}$	3.66 ± 0.25 b**
(200 ml/kg)				
Diabetic + Glibenclamide	104.15 ± 1.25 b**	3.45 ± 0.26 b**	2.70 ± 0.15 b**	3.80 ± 0.26 b**
(0.5 mg/kg)				

Each value is expressed as Mean \pm SEM, where n=6. a Normal control group vs. diabetic control group, b Diabetic control group vs. all treated group, **p < 0.001.





RESEARCH ARTICLE

In silico Analysis to Validate the Activity of Quinazolinone Derivatives against Type 2 Diabetes Mellitus - A Computational Drug Discovery Study

Mincy Mathew^{1*}, D.Kilimozhi², Santhosh M Mathews³ and Anton Smith⁴

¹Research Scholar, Department of Pharmacy, Faculty of Engineering and Technology (Affiliated to Annamalai University, Chidambaram) Annamalai Nagar, Tamil Nadu, India.

²Associate Professor, Department of Pharmacy, Faculty of Engineering and Technology, (Affiliated to Annamalai University, Chidambaram) Annamalai Nagar, Tamil Nadu, India.

³Professor, Department of Pharmaceutics, Pushpagiri College of Pharmacy, (Affiliated to Kerala University of Health Sciences) Kerala, India.

⁴Professor, Department of Pharmacy, Faculty of Engineering and Technology, (Affiliated to Annamalai University, Chidambaram) Annamalai Nagar, Tamil Nadu, India.

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*Address for Correspondence

Mincy Mathew

Research Scholar,

Department of Pharmacy,

Faculty of Engineering and Technology

(Affiliated to Annamalai University, Chidambaram)

Annamalai Nagar, Tamil Nadu, India.

Email: min_cmath@yahoo.co.in



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ABSTRACT

Type 2 Diabetes Mellitus (T2DM) is an escalating worldwide health concern, demanding the development of innovative therapeutic strategies or potential drug candidates. This research employed an insilico analysis to validate the antidiabetic activity of novel quinazolinone derivatives against multiple T2DM-associated biomolecules such as alpha-amylase, insulin-like growth factor 1 receptor, sulfonylurea receptor -1, glycogen phosphorylase, lysosomal acid-alpha-glucosidase, glycogen synthase kinase-3 beta, dipeptidyl peptidase, and peroxisome proliferator-activated receptor gamma. Data mining, molecular modeling, molecular property calculation, pharmacokinetic-pharmacodynamic studies, molecular docking, and molecular dynamic simulations are the computational approaches encompassed for the study. QZ2 is identified as a promising candidate, exhibiting low toxicity, high drug-likeness, and favorable pharmacokinetics with remarkable binding affinities with peroxisome proliferator-activated receptor gamma (4PRG: -10.8 kcal/mol) and sulfonylurea receptor-1 (6JB3: -9.2 kcal/mol). The findings offer critical insights into potential antidiabetic compounds, guiding future in vitro, and in vivo experimental investigations and further applications as therapeutics for T2DM.





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Keywords: Diabetes; Quinazolinone; Molecular modeling; Molecular docking; Molecular dynamic simulation.

INTRODUCTION

Diabetes mellitus is a metabolic disorder characterized by hyperglycemia, resulting from either insufficient insulin production by the pancreas or impaired insulin responsiveness within the body. In 2019, the statistical projection indicated that the prevalence of diabetes in India encompassed approximately 77 million individuals, and projections indicate that this figure is poised to surge, reaching an anticipated 134 million cases by the year 2045 [1]. Type 2 Diabetes Mellitus (T2DM), recognized as the ninth most prevalent cause of mortality worldwide, is linked to a spectrum of multiorgan complications, encompassing both microvascular and macrovascular issues, resulting in premature morbidity and mortality, emphasizing the urgent need for comprehensive research and interventions. Its increasing global incidence and the associated burden on healthcare systems underscore the urgent need for effective treatment strategies [2]. An effective control of blood sugar, blood pressure, and blood lipid levels can serve as a preventive measure, delaying the onset of complications associated with Diabetes. Despite their diverse and distinct functions, the biomolecules alpha-amylase, insulin-like growth factor 1 receptor, sulfonylurea receptor -1, glycogen phosphorylase, lysosomal acid-alpha-glucosidase, glycogen synthase kinase-3 beta, dipeptidyl peptidase, and peroxisome proliferator-activated receptor gamma play crucial roles in diabetes and insulin regulation, alpha-amylase plays a role in carbohydrate metabolism, affecting glucose levels [3].

Insulin-like growth factor 1 receptor is implicated in the insulin signaling pathway, impacting growth and metabolism [4]. Sulfonylurea receptor-1 is a key component in the regulation of insulin secretion [5]. Glycogen phosphorylase and glycogen synthase kinase-3 beta influence glycogen storage and release, which is crucial in glucose homeostasis [6] [7]. Lysosomal acid-alpha-glucosidase is essential for glycogen breakdown, a process linked to certain types of diabetes [8]. Dipeptidyl peptidase is involved in regulating the activity of incretin hormones, which impact insulin secretion [9]. Peroxisome proliferator-activated receptor gamma plays a role in insulin sensitivity and glucose metabolism [10]. In various ways, these biomolecules contribute to the intricate network of processes related to diabetes and insulin regulation in the human body. Many heterocyclic compounds both natural and synthetic exhibit antidiabetic activity. The fundamental structural scaffolds of such pharmacological agents encompass thiazolidinone, azole, chalcone, pyrrole, pyrimidine, and various others [11]. Diverse drug classes employ distinct mechanisms for regulating blood glucose levels. However, synthetic drugs are all associated with a plethora of adverse effects. Hence, it is essential to explore novel compounds that can more effectively combat diabetes with fewer side effects. Quinazolinone represents a fused nitrogen-containing heterocyclic ring structure. The quinazolinone moiety has been associated with a wide spectrum of pharmacological activities, including anticancer, anticonvulsant, antimalarial, antifungal, antibacterial, anti-inflammatory, antidiabetic, and hypolipidemic effects.

There are some approved drugs in the market as anticancer agents such as EGFR inhibitors, antihypertensive as alpha receptor blockers, etc. which contain quinazolinone as a core heterocyclic moiety [12]. Traditional drug discovery approaches are often time-consuming and resource-intensive, making it imperative to explore innovative methods to expedite the identification of potential therapeutics. Computational drug discovery, an indispensable tool in modern pharmaceutical research, offers an efficient and cost-effective means to assess the activity of novel molecules against target molecules. *In silico* analysis, as a key component of this approach, enables the prediction and validation of molecular interactions, thereby streamlining the drug discovery process. The study delves into the realm of computational drug discovery, to validate the activity of novel molecules against T2DM. For this, we designed various quinazolinone nucleus derivatives and performed various *in silico* analyses. By harnessing the power of computational approaches, we aim to accelerate the identification of promising drug candidates with antidiabetic activity with a reduced reliance on resource-intensive laboratory experiments.





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MATERIALS AND METHODS

The research methodology includes a series of computational approaches such as Data mining, Molecular modeling, Molecular Property Calculation, Molecular docking, and Molecular Dynamic simulation to identify the potential molecule. Background studies and data mining were performed on the disease, related targets, and therapeutic drug molecules. Targets and their standard drug molecules were identified and considered for the study.

TARGETS

Alpha-amylase, Insulin-like growth factor 1 receptor (IGF1R), sulfonylurea receptor -1 (SUR1), glycogen phosphorylase, lysosomal acid-alpha-glucosidase, glycogen synthase kinase-3 beta, dipeptidyl peptidase, and peroxisome proliferator-activated receptor gamma are the target molecules selected for the study. The 3D structures of the selected targets were obtained from the Protein Data Bank (PDB) (https://www.rcsb.org/) [13]. The selected PDBIDs of the protein molecules for the study are included in Table 1.

LIGANDS

Molecular Modeling

In this study, we employed various computational tools to assess the druggability of the proposed ligand molecules. ACD/ChemSketch was utilized to draw chemical structures and generate SMILE notation for the derivatives. A total of 11 novel ligand molecules - QZ1, QZ2, QZ3, QZ4, QZ5, QZ6, QZ7, QZ8, QZ9, QZ10, and QZ11 were modeled using the Avogadro Tool [14]. The modeled structures were optimized, validated, and used for further *in silico* analysis. Also, standard therapeutic drug molecules were used to assess and compare interactions and binding affinities with the docking results of the novel ligand molecules.

Molecular Property Calculations

Molecular property calculation plays a crucial role in drug discovery to assess and predict the pharmacokinetic and pharmacodynamic attributes of potential drug candidates. These calculations provide valuable insights into a compound's solubility, stability, bioavailability, toxicity, and interactions with biological targets, aiding in the selection and optimization of lead compounds with the highest likelihood of success in the development of safe and effective drugs. By leveraging molecular property data, we can streamline the drug discovery process, reduce the risk of failure in clinical trials, and accelerate the development of innovative treatments. Molinspiration Cheminformatics software was used to evaluate adherence to Lipinski's rule of five and drug-likeness properties [15]. Swiss ADME (http://www.swissadme.ch/) &PkCSM software (https://biosig.lab.uq.edu.au/pkcsm/) were employed to compute physicochemical properties, predict ADME parameters, assess pharmacokinetic properties, and determine drug-like qualities [16][17]. Additionally, we employed OSIRIS Property Explorer (https://openmolecules.org/propertyexplorer/) to anticipate properties associated with high risks of undesired effects, such as mutagenicity or suboptimal intestinal absorption.

Screening of Ligands

The molecular properties of ligands were screened under Lipinski's rule of five and ADMET properties. Lipinski's Rule of Five is a set of 5 rules used in drug discovery and medicinal chemistry to evaluate the drug-likeness and potential for oral bioavailability of chemical compounds [18]. The rule criteria include,

- 1. Molecular Weight: should be less than 500 daltons (Da).
- 2. Lipophilicity (LogP): should be less than 5.
- 3. Hydrogen Bond Donors: should have no more than 5 hydrogen bond donor atoms (nitrogen and oxygen).
- 4. Hydrogen Bond Acceptors: should have no more than 10 hydrogen bond acceptor atoms (nitrogen and oxygen).
- 5. Molar refractivity: should be between 40-130.





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ADMET calculations involve assessing the absorption, distribution, metabolism, excretion, and toxicity properties of a pharmaceutical compound [19]. The process plays a crucial role in drug discovery and development by helping to predict the behavior of compounds within the body.

MOLECULAR DOCKING STUDIES

The molecular interaction analysis (molecular docking) of the targets and ligand molecules was performed using Autodock Vina software (v.1.2.0.) [20]. The objective was to predict the bound conformation and the binding affinity of the considered ligands with the respective receptors. The selected targets, alpha-amylase (1B2Y), Insulin-like growth factor 1 receptor (2OJ9), sulfonylurea receptor -1 (6JB3), glycogen phosphorylase (3DD1), lysosomal acidalpha-glucosidase (5NN8), glycogen synthase kinase-3 beta (1UV5), dipeptidyl peptidase (4CDC), peroxisome proliferator-activated receptor gamma (4PRG) was docked with the modeled molecules and their corresponding standard drug molecules. The protein preparation was performed using the Swiss PDB viewer. The process includes the addition of missing residues, the addition of non-polar hydrogen, and Kollman charges. The target structure was saved in .pdbqt format. The grid box for each protein was generated by the AutoDock vina grid box generation application and molecular docking was performed. Autodock Vina will compute the binding affinity (Kcal mol-1) and generate the best 10 binding poses of each ligand with the receptor. This calculation relied on the ligand's conformation at the binding site and the Root Mean Square Deviation (RMSD) between the initial and docked structures.

Interaction Analysis

Interaction analysis of the receptor-ligand complex was performed using Discovery Studio Biovia 2017 (https://discover.3ds.com/discovery-studio-visualizer-download). Poses with the best binding affinity and H-bond interaction were selected and subjected to Molecular Dynamic Simulation to validate the interaction stability.

MOLECULAR DYNAMICS SIMULATION

Molecular dynamic simulations of the selected protein-ligand complexes were performed using the GROMACS version 2020.6 MD package [21]. The process commenced with the removal of crystal water from the protein and ligand complex. Following this, the topology files of protein and ligands were created separately by using the Charmm36 force field (Charmm36-jul 2021.ff) and CGenFf server respectively. Subsequently, the systems were solvated within a dodecahedron box utilizing the transferable intermolecular potential with a three-point (TIP3P) water model. To maintain neutrality, Na+ and Cl- ions were added. Energy minimization was conducted using the steepest descent integrator to resolve any steric conflicts between the proteins and water molecules. After energy minimization, ligand restraining, and temperature coupling were performed. The systems underwent a two-step equilibration process: primarily volume equilibration in the NVT ensemble (constant number of particles, volume, and temperature) and then pressure equilibration in the NPT ensemble (constant number of particles, system pressure, and temperature). Both equilibration steps consisted of 100,000 steps, resulting in a cumulative simulation time of 200 picoseconds (ps). A constant pressure of 1 atm and a temperature of 300 K were maintained. Following, a production molecular dynamics run was conducted, consisting of 50,000,000 steps, equivalent to 100 nanoseconds (ns).

Trajectory Plot Generation And Analysis

MD simulation result analysis encompassed the analysis of trajectory plots and receptor-ligand interactions. The trajectory plots were generated and analyzed using Xmgrace. RMSD, RMSF, RoG, H-bond interaction distribution, and SASA were analyzed to assess the stability of the interaction between the biomolecules [22]. RMSD measures the deviation or difference between the coordinates of atoms in a given structure and a reference structure. It provides a quantitative measure of the molecule's confirmation changes over time during a simulation. This is essential for assessing the stability and structural changes of biomolecules or ligand-receptor complexes. RMSF provides insight into the flexibility and stability of individual atoms or residues within a biomolecular system and helps to identify regions of significant structural variability. By understanding RMSF, we can pinpoint critical regions that undergo conformational changes, aiding in the interpretation of protein dynamics and interactions. The protein compactness





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was studied by plotting the radius of gyration (Rg). The number of Hydrogen bonds formed between the protein and ligands for 100 ns was depicted. For visualizing the post-MD simulation complexes and interaction, various tools, including DS Visualizer, Visual Molecular Dynamics, and Pymol, were employed. Protein compactness and overall shape of a biomolecule were evaluated using the radius of gyration (Rg) plot. Furthermore, the number of hydrogen bonds formed between the protein and ligands during the 100 ns period was illustrated in the H-bond distribution plot. Solvent Accessible Surface Area (SASA) plots provide insights into the structural dynamics and accessibility of specific regions, aiding in the characterization of biomolecular interactions and conformational changes. The gmx_MMPBSA package of GROMACS is used to calculate the binding free energy. CHARMM force field is used for this computational approach. Using the command "gmxmmpbsa, "the binding free energy of a complex was calculated from 200 frames extracted from the 100 ns MD trajectory.

RESULT

The *in silico* analysis was conducted to validate the anti diabetic activity of the newly modeled molecules against multiple targets that contribute to an intricate network of processes associated with diabetes and insulin regulation in the human body. The study focused on various proteins that play a role in the complex network of processes associated with diabetes and the regulation of insulin in the human body (Table 1).

Designing Of the Novel Molecules and Structure Modeling

A total of 11 quinazolinone derivatives were modeled and structures were optimized. The structure of the novel molecules is included in Table 2. These structures were modeled in 2D .sdf file format and subjected to further analysis.

Molecular Properties and ADMET Prediction.

Molecular and ADMET properties were calculated using various software and molecules were screened based on Lipinski's rule (Table 3). Table 3: The molecular descriptors include miLogP (predicted octanol-water partition coefficient), TPSA (topological polar surface area), nAtoms (number of atoms), MW (molecular weight), nON (number of oxygen atoms), nOHNH (number of hydroxyl and amine groups), nViolations (number of Lipinski rule violations), nRotb (number of rotatable bonds), and Volume. Seven of the eleven modeled molecules satisfied Lipinski's rule of five. QZ1, QZ2, QZ6, QZ7, QZ8, QZ9, and QZ11 were considered for next-level analysis, i.e., Toxicity Risk, Drug Likeness, and Solubility predictions (Table 4, 5, 6, 7). Table 4: The toxicity prediction of the modeled molecules: Toxicity Risk (Mutagenic, Tumorigenic, Irritant); Reproductive effect; Drug Likeness; Drug score; Solubility (Clog P - lipophilicity& Clog S - aqueous solubility). Out of the seven considered molecules, QZ11 shows a high tumorigenic property. As a result, the subsequent analysis was carried out using the remaining six molecules. The pharmacokinetics (ADME), and pharmacodynamic properties were calculated by using various software (Table 5& 6). Table 5: The prediction of Absorption, Blood-brain barrier (BBB) permeability, Bioavailability, Human intestinal absorption, and Caco2 permeability. Table 6: The prediction of P-glycoprotein, Cytochrome P450 activity of modeled molecules whether they acting as substrates or inhibitors. Table 7: The prediction of P-glycoprotein, Cytochrome P450 activity of modeled molecules whether they acting as substrates or inhibitors.

Molecular Docking by AutoDock Vina

The molecular interaction analysis of the human proteins and the ligand molecules were performed using Autodock Vina. Standard drug molecules for each receptor were also considered for the study, to perform comparative analysis. The dock results of the eight receptor-ligand complexes were analyzed, screened, and ranked based on the binding affinity and hydrogen bonding interactions (Table 8). Table 8: Dock result of multiple therapeutic targets-standard drug and modeled molecules. RECEPTORS (PDB ID): Alpha-amylase (1B2Y), Insulin-like growth factor 1 receptor (2OJ9), Sulfonylurea receptor -1 (6JB3), Glycogen phosphorylase (3DD1), Lysosomal acid-alpha-glucosidase (5NN8), Glycogen synthase kinase-3 beta (1UV5), Dipeptidyl peptidase (4CDC), Peroxisome proliferator-activated receptor gamma (4PRG). The results of the eight receptor-ligand docking show a binding affinity ranging from -4.8





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kcal/mol to -10.8 kcal/mol. The selected receptor-ligand complexes were considered for further analysis (Figure 1 & 2).

Molecular Dynamic Simulation

To confirm the stability of the interactions, Molecular dynamic simulations of the top-ranked ligand poses were performed to validate the interaction stability. Following the MD simulation, we analyzed the trajectory and interactions to gain insights into the protein's spatial fluctuations and changes in interactions. The outcomes encompass a spectrum of interactions, including both stable and unstable ones. Some ligands maintain interactions with the same amino acid residues as initially observed in the molecular docking results. Conversely, others exhibit different interactions, primarily influenced by fluctuations in energy and shifts in conformational geometry (Table 9). Table 9: Molecular Docking and Molecular Dynamic simulation interaction result of Peroxisome proliferator-activated receptor gamma (4PRG) and Sulfonylurea receptor -1 (6JB3). Following molecular dynamic simulation, a thorough examination of interaction and trajectory was performed (Figure 3 and Figure 4) to validate the stability of the receptor- ligand interaction. Two complexes 4PRG - QZ2 and 6JB3-QZ2 were subjected to MMPBSA calculation. The score was found to be -41.3 kcal/mol for 4PRG - QZ2 complex and -21.26 kcal/mol for 6JB3 – QZ2 complex.

DISCUSSIONS

The insilico analysis employed in this study aimed to validate the antidiabetic activity of novel quinazolinone analogs against multiple targets associated with Type 2 Diabetes Mellitus (T2DM) disease. The urgency of effective treatment strategies for T2DM is underscored by its increasing global prevalence and the associated burden on healthcare systems [23]. The study specifically focused on biomolecules involved in the intricate network of processes related to diabetes and insulin regulation. The selected targets, including alpha-amylase, insulin-like growth factor 1 receptor, sulfonylurea receptor -1, glycogen phosphorylase, lysosomal acid-alpha-glucosidase, glycogen synthase kinase-3 beta, dipeptidyl peptidase, and peroxisome proliferator-activated receptor gamma, play significant roles in various activities related to diabetes and insulin regulation. Quinazoline derivatives were modeled and considered as ligands for the study based on their potential to interact with these targets and modulate their activities. The initial screening of ligands based on Lipinski's rule of five and ADMET properties resulted in the selection of six promising novel molecules for further analysis. The miLogP values, indicative of lipophilicity, range from 3.09 (QZ2) to 6.62 (QZ10), suggesting varying degrees of hydrophobicity among the molecules [24]. The Topological Polar Surface Area (TPSA) values, associated with molecular polarity, exhibit considerable diversity, with QZ2 having the highest TPSA (85.16 Å²), emphasizing its potential for interactions with polar biological targets [25]. Notably, the number of violations against Lipinski's rule (nViolations) is generally low across all molecules, implying their adherence to drug-likeness criteria. The number of rotatable bonds (nRotb) varies, with QZ6, QZ8, QZ9, and QZ11 having a higher count, potentially influencing their conformational flexibility [26].

Molecular weights (MW) span a broad range, with QZ1 (327.39 g/mol) being the lightest and QZ5 (485.18 g/mol) the heaviest. The number of oxygen atoms (nON) and hydroxyl/amine groups (nOHNH) is relatively consistent among the molecules, indicating a structural similarity in terms of oxygen-containing functional groups. The toxicity and drug-likeness predictions from Osiris Property Explorer for the designed quinazolinone derivatives reveal a generally favorable profile. All compounds, except QZ11, demonstrate low risks in mutagenic, tumorigenic, irritant, and reproductive effect categories. QZ11 exhibits high tumorigenic risk, indicating a potential safety concern. In terms of drug-likeness, QZ2 stands out with the highest drug likeness score (4.73) and favorable physicochemical properties (Clog P, Clog S), suggesting its potential suitability as a drug candidate. Overall, most of the compounds exhibit low toxicity risks and favorable drug-likeness properties, emphasizing QZ2 as a particularly promising candidate for further consideration in antidiabetic drug development. The subsequent pharmacokinetic and pharmacodynamic evaluations were crucial in identifying potential lead compounds with favorable drug-like properties [27]. The ADMET, bioavailability, pharmacokinetic properties, and toxicity predictions reveal significant insights to the research study. Notably, QZ2 demonstrates high gastrointestinal absorption, favorable water





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solubility, and remarkable human intestinal absorption, suggesting its potential as an orally administered drug [28]. QZ2 also exhibits interactions with important targets involved in diabetes regulation. However, caution is warranted due to its prediction as a P-glycoprotein substrate. Additionally, QZ2 displays promising pharmacokinetic attributes, including steady-state volume of distribution and BBB permeability. Despite its predicted CYP3A4 inhibition and AMES toxicity, further experimental validation is crucial to confirm the translational potential of QZ2 as a potential antidiabetic drug candidate. The overall findings highlight the importance of integrating computational predictions to guide the selection of lead compounds for subsequent experimental investigations. The molecular docking studies provided insights into the binding affinities and interactions between the ligands and target proteins [29]. As a result of the interaction analysis, QZ2 stands as a promising candidate, exhibiting remarkable binding affinities and consistent stability in interactions with the selected targets, particularly Peroxisome proliferator-activated receptor gamma (4PRG) and Sulfonylurea receptor-1 (6JB3). QZ2 exhibits a binding affinity of -10.8 kcal/mol to 4PRG and -9.2 kcal/mol to 6JB3. In the 4PRG-QZ2 complex, the ligand interacts with amino acid residues GLU 343, SER 342, and MET 364. In the 6JB3-QZ2 complex, the interaction formed with ASN 1296 and ASN B 547 residues during molecular docking. The stability of these interactions was validated through Molecular Dynamics (MD) simulations over a 100 ns period [30]. Notably, in the 4PRG-QZ2 complex, the ligand maintains a stable interaction with GLU 343 and SER 342 residues, persisting even after a 100 ns MDS (Figure 2). Similarly, in the case of the 6JB3-QZ2 complex, it sustains stability with ASN 1296 throughout a 100 ns MD simulation study (Figure 3).

The analysis of trajectory plots, including RMSD, RMSF, RoG, H-bond interactions, and SASA, offered a comprehensive understanding of the ligand-protein dynamics [31]. For the 4PRG-QZ2 complex, the RMSD plot recorded a minimum value at 0.1 ns (0.134272 nm) and a maximum at 7.30 ns (0.270155 nm). The plot indicates that the protein maintains relative stability in the time from 50.9 ns to 100 ns. The protein encompasses residues from GLU 207 to LEU 476. The RMSF plot for the 4PRG-QZ2 complex reveals fluctuations ranging from 0.0389 nm to 0.5574 nm across the entire protein. Notably, Leu 476 displays the highest fluctuation at 0.5574 nm. The radius of gyration (RoG) plot for 4PRG-QZ2 suggests a deviation in Rg from 1.93337 nm (at 7.4 ns) to 1.99864 nm (at 1.9 ns), with an average of 1.962889 nm. During the 100 ns molecular dynamics (MD) simulation, a maximum of 3 hydrogen bonds were observed. The solvent-accessible surface area (SASA) plot of the 4PRG-QZ2 complex indicates structural fluctuations ranging from 143.172 nm2 (at 11.9 ns) to 158.389 nm2 (at 0.9 ns). The SASA value is 141.402 nm2 at 0 ns and 151.442 nm2 at 100 ns. In the case of the 6JB3-QZ2 complex, the RMSD values depicted in the plot exhibit a range with a minimum of 0.2364022 nm observed at 0.1 ns and a maximum of 0.50818 nm at 50.80 ns. This graphical representation suggests that the protein maintains relative stability between 56.6 ns and 100 ns. The residue count spans from VAL 215 to TYR 1326. The RMSF plot for the 6JB3-QZ2 complex reveals fluctuations across the entire protein, ranging from 0.0498 nm to 0.4765 nm. Notably, TYR 1326 experiences the highest fluctuation at 0.4765 nm.

Examining the radius of gyration (RoG) plot for 6JB3-QZ2, it is observed that the Rg deviation varies from 2.74061 nm at 26.4 ns to 2.78635 nm at 80.7 ns, averaging at 2.763852 nm. Throughout the 100 ns molecular dynamics simulation, a maximum of 2 hydrogen bonds are observed. The SASA plot for the 6JB3-QZ2 complex indicates structural fluctuations ranging from 298.411 nm² at 91.3 ns to 319.825 nm² at 41.4 ns. The SASA value at the initial time point (0 ns) is 300.78 nm², and after the simulation (100 ns), it is 300.516 nm². The results collectively highlight QZ2 as a promising candidate with potential antidiabetic activity, demonstrating stable interactions with key biomolecules associated with T2DM. However, further experimental validation is essential to confirm the in silico findings and assess the translational potential of QZ2 as a therapeutic agent for T2DM. In conclusion, this computational drug discovery study provides valuable insights into the design and evaluation of novel quinazolinone derivatives for their potential antidiabetic activity. The integration of in silico analyses, including molecular docking and MD simulations, offers a systematic approach to identifying lead compounds with the desired pharmacological properties. The findings lay the foundation for future experimental investigations and the development of innovative therapeutics for Type 2 Diabetes Mellitus.





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CONCLUSION

The research study undertook, *in silico* analysis, to validate the antidiabetic potential of quinazolinone derivatives against multiple Type 2 Diabetes Mellitus (T2DM) targets. The modeled quinazolinone derivative molecules were analyzed against key biomolecules - alpha-amylase, insulin-like growth factor 1 receptor, sulfonylurea receptor -1, glycogen phosphorylase, lysosomal acid-alpha-glucosidase, glycogen synthase kinase-3 beta, dipeptidyl peptidase, and peroxisome proliferator-activated receptor gamma, which are associated with diabetes and insulin regulation. The Computational approaches to molecular docking and dynamics identified QZ2 as a promising candidate with low toxicity, high drug-likeness, and favorable pharmacokinetic attributes. Noteworthy binding affinities were observed with Peroxisome proliferator-activated receptor gamma (4PRG, -10.8kcal/mol) and sulfonylurea receptor-1 (6JB3, -9.2 kcal/mol). These findings contribute valuable insights into the antidiabetic potentiality of the QZ2 for future *in vitro*, and *in vivo* experimental investigations and the development of innovative therapeutics for T2DM.

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Table 1: Proteins selected for the study; corresponding PDB IDs and the therapeutic drug molecules.

Sl No:	Protein Name	PDB ID	Standard Drugs
1	Alpha-amylase	1B2Y	ACARBOSE
2	Insulin-like growth factor 1 receptor (IGF1R)	2OJ9	METFORMIN
3	Sulfonylurea receptor -1 (SUR1)	6JB3	REPAGLINIDE
4	Glycogen phosphorylase	3DD1	METFORMIN





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5	Lysosomal acid-alpha-glucosidase	5NN8	ACARBOSE
6	Glycogen synthase kinase-3 beta	1UV5	AR-AO14418
7	Dipeptidyl peptidase	4CDC	SITAGLIPTIN
8	Peroxisome proliferator-activated receptor	4PRG	PIOGLITAZON
	gamma		E

Table 2: Structure of the proposed derivatives drawn by ACD/ChemSketch

S1. No:	Sample code	Proposed Derivatives	Name of the Compound
1.	QZ1		3-[(E)-benzylideneamino]- 2 -phenyl- 2 , 3 -dihydroquinazolin- 4 ($1H$)-one
2.	QZ2		2-(4-hydroxyphenyl)-3-{(<i>E</i>)-[(4-hydroxyphenyl)methylidene]amino}-2,3-dihydroquinazolin-4(1 <i>H</i>)-one
3	QZ3		2-(4-chlorophenyl)-3- $\{(E)$ - $[(4$ -chlorophenyl)methylidene]amino}-2,3-dihydroquinazolin- $4(1H)$ -one
4	QZ4		2-(3-chlorophenyl)-3- $\{(E)$ - $[(3$ -chlorophenyl)methylidene]amino}-2,3-dihydroquinazolin- $4(1H)$ -one
5	QZ5		2-(4-bromophenyl)-3-{(<i>E</i>)-[(4-bromophenyl)methylidene]amino}-2,3-dihydroquinazolin-4(1 <i>H</i>)-one
6	QZ6		2-(4-ethoxyphenyl)-3- $\{(E)$ - $[(4$ -ethoxyphenyl)methylidene]amino}-2,3-dihydroquinazolin- $4(1H)$ -one





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7	QZ7	2-(4-methylphenyl)-3- $\{(E)$ - $[(4$ -methylphenyl)methylidene]amino}-2,3-dihydroquinazolin- $4(1H)$ -one
8	QZ8	2-(2,4-dimethoxyphenyl)-3- $\{(E)$ - $[(2,4-dimethoxyphenyl)$ -3-dihydroquinazolin- $\{(1H)$ -one
9	QZ9	$2\text{-}(3,4\text{-}dimethoxyphenyl)\text{-}3\text{-}\{(E)\text{-}[(3,4\text{-}dimethoxyphenyl})\text{-}2,3\text{-}dihydroquinazolin\text{-}4}(1H)\text{-}one$
10	QZ10	2-(2,4-dichlorophenyl)-3-{(<i>E</i>)-[(2,4-dichlorophenyl)methylidene]amino}- 2,3-dihydroquinazolin-4(1 <i>H</i>)-one
11	QZ11	2-(N,N-dimethylaminophenyl)-3-{(E)-[(N,N-dimethylaminophenyl)methylidene]amino} -2,3-dihydroquinazolin-4(1 <i>H</i>)-one

Table 3: Predicted Lipinski Parameters and Molecular Properties of the Designed Molecules

Table	3. I leuicte	u Lipinsk	ı ı aranı	perties of the Designed Molecules						
Sl No:	Molecule	miLogP	TPSA	nAtoms	MW	nON	nOHNH	nViolations	nRotb	Volume
1	QZ1	4.05	44.70	25	327.39	4	1	0	3	300.28
2	QZ2	3.09	85.16	27	359.38	6	3	0	3	316.31
3	QZ3	5.41	44.70	27	396.28	4	1	1	3	327.35
4	QZ4	5.36	44.70	27	396.28	4	1	1	3	327.35
5	QZ5	5.67	44.70	27	485.18	4	1	1	3	336.05
6	QZ6	4.82	63.17	31	415.49	6	1	0	7	384.97
7	QZ7	4.95	44.70	27	355.44	4	1	0	3	333.40
8	QZ8	4.13	81.64	33	447.49	8	1	0	7	402.46
9	QZ9	3.34	81.64	33	447.49	8	1	0	7	402.46
10	QZ10	6.62	44.70	29	465.17	4	1	1	3	354.42
11	QZ11	4.25	51.18	31	413.52	6	1	0	5	392.09





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Table 4: Predicted Toxicity Risk, Drug Likeness, Solubility Data by Osiris Property Explorer

Sl. No:	Compound	Mutagenic	Tumorigenic	Irritant	Reproductive effect	Drug Likeness	Drug score	Clog P	Clog S
1	QZ1	LOW RISK	LOW RISK	LOW RISK	LOW RISK	4.59	0.84	3.7	-4.63
2	QZ2	LOW RISK	LOW RISK	LOW RISK	LOW RISK	4.73	0.86	3.01	-4.04
3	QZ6	LOW RISK	LOW RISK	LOW RISK	LOW RISK	3.54	0.71	4.37	-5.27
4	QZ7	LOW RISK	LOW RISK	LOW RISK	LOW RISK	3.12	0.75	4.39	5.32
5	QZ8	LOW RISK	LOW RISK	LOW RISK	LOW RISK	3.38	0.74	3.42	-4.7
6	QZ9	LOW RISK	LOW RISK	LOW RISK	LOW RISK	3.53	0.47	3.49	-4.7
7	QZ11	LOW RISK	HIGH RISK	LOW RISK	LOW RISK	6.37	0.76	3.42	-4.7

Table 5: Predicted Absorption, BBB permeability, Bioavailability, Human intestinal absorption, and Caco2 permeability by Swiss ADME.

S1 No:	Compound	GI Absorption	BBB Permeant	Bioavailability	HIA (% absorbed)	Caco2 Permeability (log Papp in 10-6 cm/s)
1	QZ1	HIGH	YES	0.55	100	1.4
2	QZ2	HIGH	NO	0.55	93.963	1.076
3	QZ6	HIGH	YES	0.55	99.06	1.168
4	QZ7	HIGH	YES	0.55	99.199	1.133
5	QZ8	HIGH	NO	0.55	100	1.283
6	QZ9	HIGH	NO	0.55	100	1.248

Table 6: Predicted P-glycoprotein and Cytochrome P450 activity

			-J F				J				
Sl	Sample	P-gp	P-gp I	P-gp II	CYP2D6	CYP3A4	CYP1A2	CYP2C19	CYP2C9	CYP2D6	CYP3A4
No:	Code	Substrate	Inhibitor	Inhibitor	Substrate	Substrate	Inhibitor	Inhibitor	Inhibitor	Inhibitor	Inhibitor
1	QZ1	Yes	Yes	Yes	No	Yes	Yes	Yes	Yes	No	No
2	QZ2	Yes	No	Yes	No	Yes	Yes	Yes	Yes	No	Yes
3	QZ6	No	Yes	Yes	No	Yes	Yes	Yes	Yes	No	Yes
4	QZ7	No	Yes	Yes	No	Yes	Yes	Yes	Yes	No	No
5	QZ8	No	Yes	Yes	No	Yes	Yes	Yes	Yes	No	Yes
6	QZ9	No	Yes	Yes	No	Yes	Yes	Yes	Yes	No	Yes

Table 7: Predicted total clearance and whether the molecule is a renal OCT2 substrate, AMES toxicity, human maximum tolerated dose, oral rat acute and chronic toxicity, hERG inhibition, hepatotoxicity, and skin sensitization.

SI No	Sample Code	Total Clearance (log/ml/min/kg)	Renal OCT2 Substrate	AMES Toxicity	Max. Tolerate Dose (log mg/kg/day)	hERG I Inhibitor	hERG II Inhibitor	Oral Rat Toxicity (LD50) (mol/kg)	Hepato- toxicity	Skin Sensitisation
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1	QZ1	0.707	No	Yes	0.54	No	Yes	2.243	Yes	No
2	QZ2	0.483	No	Yes	0.402	No	Yes	2.71	No	No
3	QZ6	0.799	No	No	0.67	No	Yes	2.757	No	No
4	QZ7	0.759	No	Yes	0.406	No	Yes	2.61	Yes	No
5	QZ8	0.807	No	No	0.581	No	Yes	2.841	No	No
6	QZ9	0.63	No	No	0.557	No	Yes	3.006	No	No

Table 8: Autodock Vina Dock score of various receptor-ligand complexes.

SI		Dock Score (kcal/mol) - Various Receptor Molecules (PDB ID)									
NO	Ligand	1B2Y	2OJ9	6JB3	3DD1	5NN8	1UV5	4CDC	4PR G		
1	Acarbose	-8.9	-	-	-	-8.5	-	-	-		
2	Metformin	-	-4.8	-	-5.9	-	-	i	ı		
3	Repaglinide	-	-	-8.6	-	-	-	i	ı		
4	Ar-AO14418	-	-	-	-	-	-8	i	ı		
5	Sitagliptin	-	-	-	-	-	-	-8.6	1		
6	Pioglitazone	-	-	-	-	-	-	1	-9.2		
7	QZ1	-8.4	-8.6	-8.7	-8.3	-7.4	-8.7	-7.8	-9.2		
8	QZ2	-8.4	-8.4	-9.2	-8	-7.7	-9.2	-9.2	-10.8		
9	QZ6	-7.8	-8.3	-8.4	-7.6	-7.2	-8.1	-7.1	-8.4		
10	QZ7	-8.3	-8.9	-9	-7.9	-7.8	-8.8	-9	-8.5		
11	QZ8	-7.7	-7.2	-8.1	-7.4	-6.9	-9.1	-7.6	-10.4		
12	QZ9	-8.1	-7.6	-8.7	-7.2	-7.6	-9	-7.2	-9.1		

Table 9: Molecular Docking and Molecular Dynamic Simulation Interaction Result of Receptor-ligand complexes.

S1 No:	Complex (Receptor- Ligand)	Molecular Docking H- bond Interaction (Distance in Å)	OtherInteracting residues in Molecular Docking (Type of Interaction)	Binding Affinity (kcal/mol)	Molecular Dynamic Simulation H-bond interaction
1	4PRG - QZ2	GLU 343 (2.6574), SER 342 (2.9249), MET 364 (2.7415)	LEU 228, LEU 333, ARG 288, LEU 330, ILE 341 (Pi-Alkyl)	-10.8	LEU 340, GLU 343, SER 342





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			CYS 285 (Pi-Sulfur)		
			ILE 281 (Carbon Hydrogen Bond)		
			ASN 1293, VAL 587 (Pi-Sigma)		
2	(IP2 ()71	ASN 1296 (2.5053), ASN	PRO 551 (Pi-Alkyl)	-9.2	ASN 1293, ASN
2	6JB3 - QZ2	B 547 (2.5715)	TRP 1297, PHE 591 (Pi-Pi T-	-9.2	1296
			shaped)		

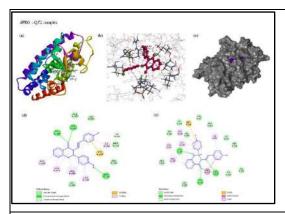


Figure 1: Molecular docking and dynamic simulation interaction diagrams of 4PRG-QZ2 complex. (a) 3D ribbon structure view of 4PRG-QZ2 interaction dock whole complex. (b) 3D line structure view of interacted fragment of 4PRG-QZ2 dock complex (pink – ligand).(c) Solid structural view of 4PRG-QZ2 docked complex. (d) 3D diagram of 4PRG-QZ2 interaction MD whole complex. (e) 3D diagram of 4PRG-QZ2 interacted fragment of MD complex. (f) 2D diagram of 4PRG-QZ2 interaction of MD complex.

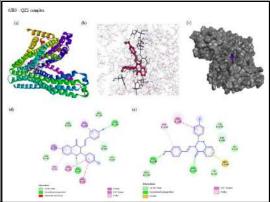


Figure 2: Molecular docking and dynamic simulation interaction diagrams of 6JB3-QZ2 complex. (a) 3D ribbon structure view of 6JB3-QZ2 interaction dock whole complex. (b) 3D line structure view of interacted fragment of 6JB3-QZ2 dock complex (pink – ligand).(c) Solid structural view of 6JB3-QZ2 docked complex. (d) 3D diagram of 6JB3-QZ2 interaction MD whole complex. (e) 3D diagram of 6JB3-QZ2 interacted fragment of MD complex. (f) 2D diagram of 6JB3-QZ2 interaction of MD complex.

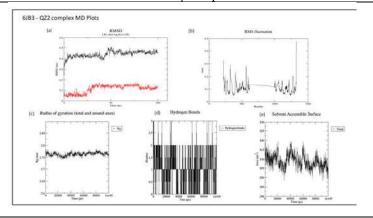


Figure 3: Trajectory plots of 4PRG-QZ2 complex (a) RMSD plot of 4PRG-QZ2 complex (b) RMSF plot of 4PRG-QZ2 complex (c) RoG plot of 4PRG-QZ2 complex (d) H-bond distribution plot of 4PRG-QZ2 complex. (e) SASA plot of 4PRG-QZ2 complex.





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RESEARCH ARTICLE

Max-Min and Min-Max Operators on Quin - Terranean Fuzzy Numbers

P.Rajarajeswari^{1*} and T.Thirunamakkani²

¹Associate Professor, Department of Mathematics, Chikkanna Government Arts College, Tirupur (Affiliated to Bharathiar University, Coimbatore) Tamil Nadu, India.

²Research Scholar, Department of Mathematics, Chikkanna Government Arts College, Tirupur (Affiliated to Bharathiar University, Coimbatore) Tamil Nadu, India.

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*Address for Correspondence

P. Rajarajeswari

Associate Professor, Department of Mathematics, Chikkanna Government Arts College, Tirupur (Affiliated to Bharathiar University, Coimbatore) Tamil Nadu, India.

Email: p.rajarajeswari29@gmail.com



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ABSTRACT

In comparison to Intuitionistic Fuzzy Set, Pythagorean Fuzzy Set and Fermatean Fuzzy Set, Quin-Terranean Fuzzy Set is a more advanced set that yields beneficial outcomes. The Quin-Terranean fuzzy set is the best measure for handling uncertainty. In this paper, we have defined some new operators namely, Max – Min $(V_{\tilde{0}})$, Min – Max $(\Lambda_{\tilde{0}})$ operators and Average operator (@) on Quin – Terranean Fuzzy Numbers. Several properties of these Max – Min (V_0) and Min – Max (Λ_0) operators as well as in combination with some basic operation already defined on Quin - Terranean Fuzzy Numbers such as complement and model operators $\boxplus \boxtimes \boxtimes \bigcirc$ are also studied. In addition, several properties of the Average operator (@) in combination with Max – Min $(V_{\bar{0}})$, Min – Max $(\Lambda_{\bar{0}})$ operators and model operators on Quin-Terranean Fuzzy numbers are also analyzed.

Keywords: Quin - Terranean fuzzy Number, Algebraic operations on Quin - Terranean fuzzy Numbers, Max - Min operator, Min - Max operator and Average operator on Quin - Terranean fuzzy Numbers.

INTRODUCTION

The idea of Fuzzy Set Theory, a development of classical set theory, was first put forward in 1965 by Zadeh [15]. For the simple computation of MIN and MAX operations for more than two upper semi-continuous Fuzzy Numbers, Dug Hun Hong and Kyung Tae Kim [2] suggested a new approach using the α -cut representation. They represented Non-continuous Fuzzy Numbers using this technique. This method allows the computation of the MIN and MAX operations for more than two Fuzzy Numbers to be done simultaneously rather than step-by-step as it is with Chiu and Wang's method. Based on Zadeh's extension principle, the max-min composition operators have a wide range of





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desired results. In 2020, for two Triangular Fuzzy Numbers defined on R, Kim Hyun *et al.* [5] determined Zadeh's max-min composition operators. To implement the decisions of a decider, in 2020, Ferdinando Di Martino and Salvatore Sessa [3] suggested a new method based on the greatest (resp., smallest) eigen fuzzy set (GEFS, resp., SEFS) of a fuzzy relation R with respect to the max-min (resp., min-max) composition. They have generated the fuzzy relations RMAX (resp., RMIN), where any entry RMAXij (resp., RMINij) expresses how much the efficacy produced on the ith characteristic is equal to or greater (resp., lesser) than that one produced by the jth characteristic. This information comes from the evaluators assessments of how much a characteristic is improved with respect to others. To increase the performances of each characteristic, the GEFS of RMAX (resp. SEFS of RMIN) are calculated. They have also suggested a method to evaluate the tourism enhancement policies in the historical centre of a significant Italian city in as a result of earlier applications based on GEFS and SEFS.

In 1986, Atanassov [1] proposed the extended notion of Fuzzy Set Theory termed as Intuitionistic Fuzzy Set (IFS) to manage instances when Fuzzy Sets are insufficient. He outlined the IFS's functions and examined their characteristics. In 2017, Ion Iancu [4] suggested a group of Intuitionistic Fuzzy similarity measures whose expressions take the hesitancy degree into consideration. This group is an expansion of a few well-known measurements. They focused in two main directions to evaluate the effectiveness of these measures: the first demonstrates the reasonableness of the suggested measures, and the second discusses the potential applications to some real-world issues. With regard to the max-min product composition operator, in 2014, Melliani *et al.* [6] expanded the LU-factorization to the Intuitionistic Fuzzy Square Matrix. They provided algorithms to find two Intuitionistic Fuzzy (lower and upper) triangular matrices L and U for an Intuitionistic Fuzzy Square Matrix A such that A = LU and a result is provided for the existence and uniqueness of this decomposition. They have also suggested a method using LU-factorization to find the square system of solutions to Intuitionistic Fuzzy Relation Equations (IFRE). They have also provided an example to highlight their work.

Yager [12] presented the Pythagorean Fuzzy Set (PFS) in 2014, an extension of the IFS in which the sum of the squares representing the acceptance and non-acceptance of a proposition is less than or equal to one. In 2022, Snehaa and Sandhiya [11] studied the idea of Pythagorean Fuzzy Sets, along with some definitions the score and accuracy function of Fuzzy sets. They analyzed few Pythagorean Fuzzy Set properties. They have established the notion of a relation known as Pythagorean Fuzzy Relation, which is supported by numerical examples. They have also proposed a career placement decision-making process based on the Pythagorean Fuzzy Relation known as min-maxmin composition which can be used to assess the appropriate suitability of careers for applicants based on skills and academic performance.

Senapathi and Yager [13] generated the Fermatean Fuzzy Set (FFS) in 2019, which is a Fuzzy Set where the sum of the cubes representing the acceptance and non-acceptance degrees is less than or equal to one. In 2019, Three additional operations—subtraction, division, and Fermatean arithmetic mean operations over Fermatean Fuzzy Numbers—were introduced by Tapan Senapati and Ronald Yager [14]. They have given a lot of information on their properties. To solve the problem of multi-criteria decision-making, they created a Fermatean Fuzzy Weighted Product Model. To support the theory they created and to demonstrate its applicability, they also offered a real-world example of selecting. In comparison to IFS, PFS, and FFS, the Quin-Terranean Fuzzy Set (QTFS) [8], an innovative concept we proposed in 2023, is the most significant and practical method to address uncertainty and contingency. Fundamental set theory operations including complement, intersection, and union were developed for Quin-Terranean Fuzzy sets. Regarding these activities, a number of the qualities are explored. The De Morgan's rule, associative property, commutative property, and distributive property have all been demonstrated to be satisfied by the Quin-Terranean Fuzzy set.

Preliminaries

In this session, the basic definitions, operations and properties related to the paper are given.





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Definition 2.1[13]

The Pythagorean fuzzy sets defined on a non-empty set X as objects having the form $\mathcal{P} = \{(x, \alpha_P(x), \beta_P(x)) : x \in X\}$, where the function $\alpha_P(x) : X \to [0,1]$ and $\beta_P(x) : X \to [0,1]$, denote the degree of membership and degree of non-membership of each element $x \in X$ to the set \mathcal{P} respectively, and $0 \le (\alpha_P(x))^2 + (\beta_P(x))^2 \le 1$ for all $x \in \mathcal{X}$. For any

Pythagorean fuzzy set \mathcal{P} and $x \in X$, $\pi_P(x) = \sqrt{1 - (\alpha_P(x))^2 - (\beta_P(x))^2}$ is called the degree of indeterminacy of x to \mathcal{P} .

Definition 2.2[13]

Let X be a universe of discourse. A Fermatean fuzzy set \mathcal{F} in X is an object having the form

 $\mathcal{F} = \{ \langle x, \alpha_{\mathcal{F}}(x), \beta_{\mathcal{F}}(x) \rangle : x \in X \},$

where $\alpha_{\mathcal{F}}(x): X \to [0,1]$ and $\beta_{\mathcal{F}}(x): X \to [0,1]$, including the condition

$$0 \le \left(\alpha_P(x)\right)^3 + \left(\beta_P(x)\right)^3 \le 1,$$

for all $x \in X$. The $\alpha_{\mathcal{F}}(x)$ and $\beta_{\mathcal{F}}(x)$ denote, respectively, the degree of membership and the degree of non-membership of the element x in the set \mathcal{F} .

For any FFS
$$\mathcal{F}$$
 and $x \in X$,

$$\pi_{\mathcal{F}}(x) = \sqrt[3]{1 - \left(\alpha_P(x)\right)^3 - \left(\beta_P(x)\right)^3}$$

is identified as the degree of indeterminacy of x to \mathcal{F} .

Definition 2.3 [8]

Let X denotes a universe of discourse. A Quin – Terranean fuzzy set \widetilde{Q} in X is the element possess the configuration $\widetilde{Q} = \{(\widetilde{x}, \sigma_{\widetilde{O}}(\widetilde{x}), \delta_{\widetilde{O}}(\widetilde{x})): \forall \widetilde{x} \in X\}$,

where $\sigma_{\widetilde{Q}}(\widetilde{x}): X \to [0,1]$ and $\delta_{\widetilde{Q}}(\widetilde{x}): X \to [0,1]$ represents the degree of dependence and degree of non – dependence of the element respectively, satisfying the condition $0 \le \left(\sigma_{\widetilde{Q}}(\widetilde{x})\right)^{4^5} + \left(\delta_{\widetilde{Q}}(\widetilde{x})\right)^{\frac{5}{4}} \le 1, \forall \widetilde{x} \in X$.

For any Quin – Terranean fuzzy set \widetilde{Q} and $\forall \widetilde{x} \in X$, $\varphi_{\widetilde{Q}}(\widetilde{x}) = \left\{ 1 - \left(\sigma_{\widetilde{Q}}(\widetilde{x}) \right)^{\frac{4}{5}} - \left(\delta_{\widetilde{Q}}(\widetilde{x}) \right)^{\frac{5}{4}} \right\}^{\frac{4}{5}}$ is the degree of hesitance

of $\widetilde{\boldsymbol{x}}$ to \boldsymbol{X} .

Definition 2.4 [7] A Quin – Terranean fuzzy number \tilde{Q} is defined as,

- (a) a Quin Terranean fuzzy subset of the real line
- (b) normal, that is, there exist $\tilde{x}_0 \in R$ such that $\sigma_{\tilde{o}}(\tilde{x}_0) = 1$ and $\delta_{\tilde{o}}(\tilde{x}_0) = 0$.
- (c) a convex set for the acceptance function $\sigma_{\tilde{o}}(\tilde{x})$, that is,

$$\sigma_{\tilde{\mathcal{O}}}(\lambda \tilde{x}_1 + (1 - \lambda)\tilde{x}_2) \ge \min\left(\sigma_{\tilde{\mathcal{O}}}(\tilde{x}_1), \sigma_{\tilde{\mathcal{O}}}(\tilde{x}_2)\right) \qquad \forall \, \tilde{x}_1, \tilde{x}_2 \in R, \lambda \in [0, 1]$$

(d) a concave set for the non-acceptance function $\delta_{\tilde{o}}(\tilde{x})$, that is,

$$\delta_{\widetilde{Q}}(\lambda \widetilde{x}_1 + (1 - \lambda)\widetilde{x}_2) \leq \max \left(\delta_{\widetilde{Q}}(\widetilde{x}_1), \delta_{\widetilde{Q}}(\widetilde{x}_2)\right) \qquad \forall \ \widetilde{x}_1, \widetilde{x}_2 \in R, \lambda \in [0, 1].$$

and satisfies the condition, $0 \le \left(\sigma_{\widetilde{Q}}\left(\widetilde{x}\right)\right)^{4^{5}} + \left(\delta_{\widetilde{Q}}\left(\widetilde{x}\right)\right)^{\frac{2}{4}} \le 1$.

A doublet expressed by $\widetilde{\mathbf{Q}} = \langle \sigma_{\widetilde{\mathbf{Q}}}(\widetilde{x}), \delta_{\widetilde{\mathbf{Q}}}(\widetilde{x}) \rangle$ is known as Quin – Terranean fuzzy number. The Zero and one Quin – Terranean fuzzy number is denoted by (0,0) and (1,0) respectively.

Definition 2.5 [9]

Let $\widetilde{\mathbb{Q}} = \langle \sigma_{\widetilde{\mathbb{Q}}}(\widetilde{x}), \delta_{\widetilde{\mathbb{Q}}}(x) \rangle$, $\widetilde{\mathbb{Q}}_1 = \langle \sigma_{\widetilde{\mathbb{Q}}_1}(\widetilde{x}), \delta_{\widetilde{\mathbb{Q}}_1}(x) \rangle$ and $\widetilde{\mathbb{Q}}_2 = \langle \sigma_{\widetilde{\mathbb{Q}}_2}(\widetilde{x}), \delta_{\widetilde{\mathbb{Q}}_2}(\widetilde{x}) \rangle$ be three Quin – Terranean Fuzzy Numbers and $\lambda > 0$, then their operations are expressed as follows:

(i)
$$\widetilde{\mathbb{Q}}_1 \coprod \widetilde{\mathbb{Q}}_2 = \langle \left\{ \left(\sigma_{\widetilde{\mathbb{Q}}_1}(\widetilde{x}) \right)^{4^5} + \left(\sigma_{\widetilde{\mathbb{Q}}_2}(\widetilde{x}) \right)^{4^5} - \left(\sigma_{\widetilde{\mathbb{Q}}_1}(\widetilde{x}) \right)^{4^5} \left(\sigma_{\widetilde{\mathbb{Q}}_2}(\widetilde{x}) \right)^{4^5} \right\}^{\frac{1}{4^5}}, \delta_{\widetilde{\mathbb{Q}}_1}(\widetilde{x}) \delta_{\widetilde{\mathbb{Q}}_2}(\widetilde{x}) \rangle$$

$$(ii) \qquad \widetilde{\mathbb{Q}}_1 \boxtimes \widetilde{\mathbb{Q}}_2 = \langle \, \sigma_{\widetilde{\mathbb{Q}}_1}(\widetilde{x}) \sigma_{\widetilde{\mathbb{Q}}_2}(\widetilde{x}), \left\{ \left(\delta_{\widetilde{\mathbb{Q}}_1}(\widetilde{x}) \right)^{\frac{5}{4}} + \left(\delta_{\widetilde{\mathbb{Q}}_2}(\widetilde{x}) \right)^{\frac{5}{4}} - \left(\delta_{\widetilde{\mathbb{Q}}_1}(\widetilde{x}) \right)^{\frac{5}{4}} \left(\delta_{\widetilde{\mathbb{Q}}_2}(\widetilde{x}) \right)^{\frac{5}{4}} \right\}^{\frac{1}{5}}$$





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$$(\mathrm{iii}) \qquad \lambda \widetilde{\mathbb{Q}} = \langle \left\{ 1 - \left(1 - \left(\sigma_{\widetilde{\mathbb{Q}}}(\widetilde{x}) \right)^{4^5} \right)^{\lambda} \right\}^{\frac{1}{4^5}}, \delta_{\widetilde{\mathbb{Q}}}(\widetilde{x})^{\lambda} \rangle$$

$$(\mathrm{iv}) \qquad \widetilde{\mathbb{Q}}^{\lambda} = \langle \sigma_{\widetilde{\mathbb{Q}}}(\widetilde{x})^{\lambda}, \left\{ 1 - \left(1 - \left(\delta_{\widetilde{\mathbb{Q}}}(\widetilde{x}) \right)^{\frac{5}{4}} \right)^{\lambda} \right\}^{\frac{4}{5}} \rangle.$$

Definition 2.6 [10]

Let $T_1 = \langle \sigma_{T_1}, \delta_{T_1} \rangle$ and $T_2 = \langle \sigma_{T_2}, \delta_{T_2} \rangle$ be two Quin-Terranean Fuzzy numbers, then,

(i)
$$T_1 \ominus T_2 = \langle \left\{ \frac{\sigma_{T_1}^{4^5} - \sigma_{T_2}^{4^5}}{1 - \sigma_{T_2}^{4^5}} \right\}^{\frac{1}{4^5}}, \frac{\delta_{T_1}}{\delta_{T_2}} \rangle, \text{ if } \sigma_{T_1} \geq \sigma_{T_2}, \ \delta_{T_1} \leq \min \left\{ \delta_{T_2}, \frac{\delta_{T_2} \varphi_1}{\varphi_2} \right\};$$

(ii)
$$T_1 \oslash T_2 = \langle \frac{\sigma_{T_1}}{\sigma_{T_2}}, \left\{ \frac{\delta_{T_1}^{\frac{5}{4} - \delta_{T_2} \frac{5}{4}}}{1 - \delta_{T_2}^{\frac{5}{4}}} \right\}^{\frac{1}{5}} \rangle \text{, if } \sigma_{T_1} \leq min \left\{ \sigma_{T_2}, \frac{\sigma_{T_2} \varphi_1}{\varphi_2} \right\} \text{, } \delta_{T_1} \geq \delta_{T_2}.$$

MAIN RESULT

In this section, we define $\operatorname{Max} - \operatorname{Min}(V_{\widehat{Q}})$, $\operatorname{Min} - \operatorname{Max}(\Lambda_{\widehat{Q}})$ operators and Average operator (@) on Quin – Terranean Fuzzy Numbers. Several properties of these $\operatorname{Max} - \operatorname{Min}(V_{\widehat{Q}})$ and $\operatorname{Min} - \operatorname{Max}(\Lambda_{\widehat{Q}})$ operators as well as in combination with the basic operation of complement and model operators such as $\boxplus, \boxtimes, \ominus, \oslash$ on Quin – Terranean Fuzzy Numbers are also studied. In addition, several properties of the Average operator (@) in combination with $\operatorname{Max} - \operatorname{Min}(V_{\widehat{Q}})$, $\operatorname{Min} - \operatorname{Max}(\Lambda_{\widehat{Q}})$ operators and model operators on Quin-Terranean Fuzzy numbers are also analyzed.

Max –min and Min – max operations on Quin Terranean fuzzy numbers Definition 3.1.1

Let $\tilde{L} = \langle \sigma_{\tilde{L}}(x), \delta_{\tilde{L}}(x) \rangle$, $\tilde{M} = \langle \sigma_{\tilde{M}}(x), \delta_{\tilde{M}}(x) \rangle$ be two Quin – Terranean fuzzy numbers, then the max – min $(V_{\tilde{Q}})$ and min – max $(\Lambda_{\tilde{Q}})$ operation on Quin – Terranean fuzzy numbers are defined as,

$$\widetilde{L}\bigvee_{\widetilde{Q}}\widetilde{M} = \langle max(\sigma_{\widetilde{L}}(x), \sigma_{\widetilde{M}}(x)), min(\delta_{\widetilde{L}}(x), \delta_{\widetilde{M}}(x)) \rangle$$
 and $\widetilde{L}\bigwedge_{\widetilde{Q}}\widetilde{M} = \langle min(\sigma_{\widetilde{L}}(x), \sigma_{\widetilde{M}}(x)), max(\delta_{\widetilde{L}}(x), \delta_{\widetilde{M}}(x)) \rangle$.

Properties of max –min $(\bigvee_{\tilde{O}})$ and min – max $(\bigwedge_{\tilde{O}})$ operations on Quin Terranean fuzzy numbers

The several properties of the max -min and min - max operations on Quin Terranean fuzzy numbers are given as follows,

Theorem 3.2.1: Let $\tilde{L} = \langle \sigma_{\tilde{L}}(x), \delta_{\tilde{L}}(x) \rangle$, $\tilde{M} = \langle \sigma_{\tilde{M}}(x), \delta_{\tilde{M}}(x) \rangle$ be two Quin – Terranean fuzzy numbers, then the following properties holds,

Idempotent Property:

- (i) $\tilde{L} \bigvee_{\tilde{Q}} \tilde{L} = \tilde{L}$
- (ii) $\tilde{L} \wedge_{\tilde{Q}} \tilde{L} = \tilde{L}$

Commutative property:

- (iii) $\tilde{L} \vee_{\tilde{O}} \tilde{M} = \tilde{M} \vee_{\tilde{O}} \tilde{L}$
- (iv) $\tilde{L} \wedge_{\tilde{O}} \tilde{M} = \tilde{M} \wedge_{\tilde{O}} \tilde{L}$

Demorgan Rule:

- $(v) \left(\tilde{L} \vee_{\widetilde{O}} \widetilde{M} \right)^{c} = \left(\tilde{L} \right)^{c} \wedge_{\widetilde{O}} \left(\widetilde{M} \right)^{c}$
- (vi) $(\tilde{L} \wedge_{\widetilde{O}} \tilde{M})^c = (\tilde{L})^c \vee_{\widetilde{O}} (\tilde{M})^c$

Proof:

- (i) Proof follows from the definition 3.11 (iii)
- (ii) Proof follows from the definition 3.11 (iv)
- (iii) L.H.S = $\tilde{L} \vee_{\tilde{O}} \tilde{M} = \langle max(\sigma_{\tilde{L}}(x), \sigma_{\tilde{M}}(x)), min(\delta_{\tilde{L}}(x), \delta_{\tilde{M}}(x)) \rangle$
- $= \langle \max(\sigma_{\widetilde{M}}(x), \sigma_{\widetilde{L}}(x)), \min(\delta_{\widetilde{M}}(x), \delta_{\widetilde{L}}(x)) \rangle = \widetilde{M} \vee_{\widetilde{Q}} \widetilde{L} = R. H. S.$
- (iv) L.H.S = $\tilde{L} \wedge_{\tilde{O}} \tilde{M} = \langle \min(\sigma_{\tilde{L}}(x), \sigma_{\tilde{M}}(x)), \max(\delta_{\tilde{L}}(x), \delta_{\tilde{M}}(x)) \rangle$





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= \langle \min(\sigma_{\widetilde{M}}(x), \sigma_{\widetilde{L}}(x)), \max(\delta_{\widetilde{M}}(x), \delta_{\widetilde{L}}(x)) \rangle = \widetilde{M} \wedge_{\widetilde{Q}} \widetilde{L} = R. H. S.
(v) L.H. S = (\tilde{L} \vee_{\tilde{O}} \tilde{M})^c = \langle max(\sigma_{\tilde{L}}(x), \sigma_{\tilde{M}}(x)), min(\delta_{\tilde{L}}(x), \delta_{\tilde{M}}(x)) \rangle^c
= \langle \min(\delta_{\tilde{L}}(x), \delta_{\tilde{M}}(x)), \max(\sigma_{\tilde{L}}(x), \sigma_{\tilde{M}}(x)) \rangle
=\langle \boldsymbol{\delta}_{\tilde{L}}(x), \boldsymbol{\sigma}_{\tilde{L}}(x)\rangle \wedge_{\widetilde{Q}}\langle \boldsymbol{\delta}_{\widetilde{M}}(x), \boldsymbol{\sigma}_{\widetilde{M}}(x)\rangle
= (\tilde{L})^c \wedge_{\tilde{O}} (\tilde{M})^c = R.H.S
(vi) L.H. S = (\tilde{L} \wedge_{\tilde{O}} \tilde{M})^c = \langle min(\sigma_{\tilde{L}}(x), \sigma_{\tilde{M}}(x)), max(\delta_{\tilde{L}}(x), \delta_{\tilde{M}}(x)) \rangle^c
                                                                                                                  = \langle \max(\delta_{\tilde{L}}(x), \delta_{\tilde{M}}(x)), \min(\sigma_{\tilde{L}}(x), \sigma_{\tilde{M}}(x)) \rangle
 = \langle \delta_{\tilde{L}}(x), \sigma_{\tilde{L}}(x) \rangle \vee_{\tilde{O}} \langle \delta_{\tilde{M}}(x), \sigma_{\tilde{M}}(x) \rangle = (\tilde{L})^{c} \vee_{\tilde{O}} (\tilde{M})^{c}
 = R.H.S
Theorem 3.2.2: Let \tilde{L} = \langle \sigma_{\tilde{L}}(x), \delta_{\tilde{L}}(x) \rangle, \tilde{M} = \langle \sigma_{\tilde{M}}(x), \delta_{\tilde{M}}(x) \rangle and \tilde{R} = \langle \sigma_{\tilde{R}}(x), \delta_{\tilde{R}}(x) \rangle be three Quin – Terranean fuzzy
numbers, then we have,
Associative Property:
                           \widetilde{L} \vee_{\widetilde{O}} (\widetilde{M} \vee_{\widetilde{O}} \widetilde{R}) = (\widetilde{L} \vee_{\widetilde{O}} \widetilde{M}) \vee_{\widetilde{O}} \widetilde{R}
(ii) \tilde{L} \wedge_{\tilde{O}} (\tilde{M} \wedge_{\tilde{O}} \tilde{R}) = (\tilde{L} \wedge_{\tilde{O}} \tilde{M}) \wedge_{\tilde{O}} \tilde{R}
Distributive Property:
(iii) \widetilde{L} \wedge_{\widetilde{O}} (\widetilde{M} \vee_{\widetilde{O}} \widetilde{R}) = (\widetilde{L} \wedge_{\widetilde{O}} \widetilde{M}) \vee_{\widetilde{O}} (\widetilde{L} \wedge_{\widetilde{O}} \widetilde{R})
(iv) \widetilde{L} \vee_{\widetilde{O}} (\widetilde{M} \wedge_{\widetilde{O}} \widetilde{R}) = (\widetilde{L} \vee_{\widetilde{O}} \widetilde{M}) \wedge_{\widetilde{O}} (\widetilde{L} \vee_{\widetilde{O}} \widetilde{R})
Proof:
                  L.H.S = \widetilde{L} \vee_{\widetilde{O}} (\widetilde{M} \vee_{\widetilde{O}} \widetilde{R})
   = \langle \sigma_{\tilde{L}}(x), \delta_{\tilde{L}}(x) \rangle \vee_{\tilde{O}} \langle \max(\sigma_{\tilde{M}}(x), \sigma_{\tilde{R}}(x)), \min(\delta_{\tilde{M}}(x), \delta_{\tilde{R}}(x)) \rangle
 = \langle max \left( \sigma_{\widetilde{L}}(x), max \left( \sigma_{\widetilde{M}}(x), \sigma_{\widetilde{R}}(x) \right) \right), min \left( \delta_{\widetilde{L}}(x), min \left( \delta_{\widetilde{M}}(x), \delta_{\widetilde{R}}(x) \right) \right) \rangle
 = \langle max(\sigma_{\tilde{L}}(x), \sigma_{\tilde{M}}(x), \sigma_{\tilde{R}}(x)), min(\delta_{\tilde{L}}(x), \delta_{\tilde{M}}(x), \delta_{\tilde{R}}(x)) \rangle
 = \langle max(max(\sigma_{\widetilde{L}}(x), \sigma_{\widetilde{M}}(x)), \sigma_{\widetilde{R}}(x)), min(min(\delta_{\widetilde{L}}(x), \delta_{\widetilde{M}}(x)), \delta_{\widetilde{R}}(x)) \rangle
 = \langle max(\sigma_{\tilde{L}}(x), \sigma_{\tilde{M}}(x)), min(\delta_{\tilde{L}}(x), \delta_{\tilde{M}}(x)) \rangle
          \bigvee_{\widetilde{O}} \langle \sigma_{\widetilde{R}}(x), \delta_{\widetilde{R}}(x) \rangle = (\widetilde{L} \bigvee_{\widetilde{O}} \widetilde{M}) \bigvee_{\widetilde{O}} \widetilde{R} = R.H.S.
               L.H.S = \widetilde{L} \wedge_{\widetilde{O}} (\widetilde{M} \wedge_{\widetilde{O}} \widetilde{R})
   = \langle \sigma_{\tilde{L}}(x), \delta_{\tilde{L}}(x) \rangle \wedge_{\tilde{O}} \langle x, \min(\sigma_{\tilde{M}}(x), \sigma_{\tilde{R}}(x)), \max(\delta_{\tilde{M}}(x), \delta_{\tilde{R}}(x)) \rangle
   = \langle \min \left( \sigma_{\tilde{L}}(x), \min \left( \sigma_{\tilde{M}}(x), \sigma_{\tilde{R}}(x) \right) \right), \max \left( \delta_{\tilde{L}}(x), \max \left( \delta_{\tilde{M}}(x), \delta_{\tilde{R}}(x) \right) \right) \rangle
     = \langle \min(\sigma_{\tilde{L}}(x), \sigma_{\tilde{M}}(x), \sigma_{\tilde{R}}(x)), \max(\delta_{\tilde{L}}(x), \delta_{\tilde{M}}(x), \delta_{\tilde{R}}(x)) \rangle
= \langle \min(\min(\sigma_{\tilde{L}}(x), \sigma_{\tilde{M}}(x)), \sigma_{\tilde{R}}(x)), \max(\max(\delta_{\tilde{L}}(x), \delta_{\tilde{M}}(x)), \delta_{\tilde{R}}(x)) \rangle
   = \langle \min(\sigma_{\tilde{L}}(x), \sigma_{\tilde{M}}(x)), \max(\delta_{\tilde{L}}(x), \delta_{\tilde{M}}(x)) \rangle
                  \wedge_{\widetilde{O}}\langle \sigma_{\widetilde{R}}(x), \delta_{\widetilde{R}}(x)\rangle = (\widetilde{L}\wedge_{\widetilde{O}}\widetilde{M})\wedge_{\widetilde{O}}\widetilde{R} = R.H.S.
(iii) L.H.S = \tilde{L} \wedge_{\tilde{O}} (\tilde{M} \vee_{\tilde{O}} \tilde{R})
= \langle \sigma_{\widetilde{L}}(x), \delta_{\widetilde{L}}(x) \rangle \wedge_{\widetilde{O}} \langle max(\sigma_{\widetilde{M}}(x), \sigma_{\widetilde{R}}(x)), min(\delta_{\widetilde{M}}(x), \delta_{\widetilde{R}}(x)) \rangle
= \langle \min \left( \sigma_{\tilde{L}}(x), \max \left( \sigma_{\tilde{M}}(x), \sigma_{\tilde{R}}(x) \right) \right), \max \left( \delta_{\tilde{L}}(x), \min \left( \delta_{\tilde{M}}(x), \delta_{\tilde{R}}(x) \right) \right) \rangle
       and \tilde{L} \wedge_{\tilde{O}} \tilde{M} = \langle min(\sigma_{\tilde{L}}(x), \sigma_{\tilde{M}}(x)), max(\delta_{\tilde{L}}(x), \delta_{\tilde{M}}(x)) \rangle,
              \tilde{L} \wedge_{\tilde{O}} \tilde{R} = \langle \min(\sigma_{\tilde{L}}(x), \sigma_{\tilde{R}}(x)), \max(\delta_{\tilde{L}}(x), \delta_{\tilde{R}}(x)) \rangle
R.H.S = (\tilde{L} \wedge_{\tilde{O}} \tilde{M}) \vee_{\tilde{O}} (\tilde{L} \wedge_{\tilde{O}} \tilde{R})
               \max\Big(\min\big(\sigma_{\tilde{L}}(x),\sigma_{\tilde{M}}(x)\big),\min\big(\sigma_{\tilde{L}}(x),\sigma_{\tilde{R}}(x)\big)\Big),
               min(max(\delta_{\tilde{L}}(x),\delta_{\tilde{M}}(x)), max(\delta_{\tilde{L}}(x),\delta_{\tilde{R}}(x)))
= \langle \min \left( \sigma_{\tilde{L}}(x), \max \left( \sigma_{\tilde{M}}(x), \sigma_{\tilde{R}}(x) \right) \right), \max \left( \delta_{\tilde{L}}(x), \min \left( \delta_{\tilde{M}}(x), \delta_{\tilde{R}}(x) \right) \right) \rangle
     Hence L.H.S = R.H.S, therefore, \tilde{L} \wedge_{\tilde{O}} (\tilde{M} \vee_{\tilde{O}} \tilde{R}) = (\tilde{L} \wedge_{\tilde{O}} \tilde{M}) \vee_{\tilde{O}} (\tilde{L} \wedge_{\tilde{O}} \tilde{R}).
(iv) L.H.S = \tilde{L} \vee_{\tilde{O}} (\tilde{M} \wedge_{\tilde{O}} \tilde{R})
 =\langle \, \sigma_{\tilde{L}}(x), \delta_{\tilde{L}}(x) \rangle \, \vee_{\widetilde{Q}} \, \langle \min \bigl( \sigma_{\widetilde{M}}(x), \sigma_{\widetilde{R}}(x) \bigr), \max \bigl( \delta_{\widetilde{M}}(x), \delta_{\widetilde{R}}(x) \bigr) \rangle
 = \langle max \left( \sigma_{\tilde{L}}(x), min \left( \sigma_{\tilde{M}}(x), \sigma_{\tilde{R}}(x) \right) \right), min \left( \delta_{\tilde{L}}(x), max \left( \delta_{\tilde{M}}(x), \delta_{\tilde{R}}(x) \right) \right) \rangle
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$$\begin{aligned} & \text{and } I \bigvee_{Q} \tilde{\mathbf{R}} &= \langle \max(\sigma_{1}(x), \sigma_{M}(x)), \min(\delta_{L}(x), \delta_{M}(x)) \rangle, \\ & I \bigvee_{Q} \tilde{\mathbf{R}} &\geq \langle \max(\sigma_{L}(x), \sigma_{R}(x)), \min(\delta_{L}(x), \delta_{R}(x)) \rangle, \\ & \text{RHS} &= \langle I \bigvee_{Q} \tilde{\mathbf{M}} \rangle_{Q} \langle I \bigvee_{Q} \tilde{\mathbf{R}} \rangle, \\ &= \langle \min(\max(\sigma_{L}(x), \sigma_{R}(x)), \min(\delta_{L}(x), \delta_{R}(x)) \rangle, \\ &= \langle \max(\min(\sigma_{L}(x), \delta_{R}(x)), \min(\delta_{L}(x), \delta_{R}(x)) \rangle, \\ &= \langle \max(\min(\sigma_{L}(x), \delta_{R}(x)), \min(\delta_{L}(x), \delta_{R}(x)) \rangle, \\ &= \langle \max(\min(\sigma_{L}(x), \delta_{R}(x)), \min(\delta_{L}(x), \delta_{R}(x)) \rangle, \\ &= \langle \max(\sigma_{L}(x), \min(\sigma_{R}(x), \delta_{R}(x)) \rangle, \\ &= \langle \max(\sigma_{L}(x), \delta_{L}(x)), \\ &= \langle \max(\sigma_{L}(x), \delta_{L}(x)), \\ &= \langle \max(\sigma_{L}(x), \delta_{L}(x)), \\ &= \langle \min(\sigma_{L}(x), \delta_{L}(x), \delta_{L}(x), \delta_{L}(x)), \\ &= \langle \min(\sigma_{L}(x), \delta_{L}(x), \delta_{L}(x), \delta_{L}(x)), \\ &= \langle \min(\sigma_{L}(x), \delta_{L}(x), \delta_{L}(x), \delta_{L}(x), \delta_{L}(x), \delta_{L}(x), \delta_{L}(x), \delta_{L}(x)), \\ &= \langle \min(\sigma_{L}(x), \delta_{L}(x), \delta_{$$





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 $\langle max \Big\{ \big\{ (\sigma_{\tilde{L}}(\tilde{x}))^{4^5} + (\sigma_{\tilde{R}}(\tilde{x}))^{4^5} - (\sigma_{\tilde{L}}(\tilde{x}))^{4^5} (\sigma_{\tilde{R}}(\tilde{x}))^{4^5} \big\}^{\frac{1}{4^5}}, \big\{ (\sigma_{\tilde{M}}(\tilde{x}))^{4^5} + (\sigma_{\tilde{R}}(\tilde{x}))^{4^5} - (\sigma_{\tilde{R}}(\tilde{x}))^{4^5} \big\}^{\frac{1}{4^5}}, \big\{ (\sigma_{\tilde{M}}(\tilde{x}))^{4^5} + (\sigma_{\tilde{R}}(\tilde{x}))^{4^5} - (\sigma_{\tilde{R}}(\tilde{x}))^{4^5} \big\}^{\frac{1}{4^5}}, \big\{ (\sigma_{\tilde{M}}(\tilde{x}))^{4^5} + (\sigma_{\tilde{R}}(\tilde{x}))^{4^5} - (\sigma_{\tilde{R}}(\tilde{x}))^{4^5} + (\sigma_{\tilde{R}}(\tilde{x}))^{4^5} + (\sigma_{\tilde{R}}(\tilde{x}))^{4^5} - (\sigma_{\tilde{R}}(\tilde{x}))^{4^5} + (\sigma_{\tilde{R}$ $(\sigma_{\widetilde{M}}(\widetilde{x}))^{4^5}(\sigma_{R}(\widetilde{x}))^{4^5}\}^{\frac{1}{4^5}}, min\{\delta_{L}(\widetilde{x})\delta_{R}(\widetilde{x}), \delta_{\widetilde{M}}(\widetilde{x})\delta_{R}(\widetilde{x})\}$ $= \langle max \bigg\{ \Big\{ \Big(1 - (\sigma_{\tilde{R}}(\tilde{x}))^{4^5} \Big) (\sigma_{\tilde{L}}(\tilde{x}))^{4^5} + (\sigma_{\tilde{R}}(\tilde{x}))^{4^5} \Big\}^{\frac{1}{4^5}}, \Big\{ \Big(1 - (\sigma_{\tilde{R}}(\tilde{x}))^{4^5} \Big) (\sigma_{\tilde{M}}(\tilde{x}))^{4^5} \Big\}^{\frac{1}{4^5}} \Big\} \bigg\}$ $+(\sigma_{\mathcal{B}}(\tilde{x}))^{4^{5}}\}^{\frac{1}{4^{5}}}_{4^{5}}, min\{\delta_{\tilde{I}}(\tilde{x})\delta_{\mathcal{B}}(\tilde{x}),\delta_{\tilde{M}}(\tilde{x})\delta_{\mathcal{B}}(\tilde{x})\}$ $=\langle\{\left(1-(\sigma_{\tilde{R}}(\tilde{x}))^{4^{5}}\right)max\{\sigma_{\tilde{L}}(\tilde{x})^{4^{5}},\sigma_{\tilde{M}}(\tilde{x})^{4^{5}}\}+(\sigma_{\tilde{R}}(\tilde{x}))^{4^{5}}\}^{\frac{1}{4^{5}}},min\{\delta_{\tilde{L}}(\tilde{x})\delta_{\tilde{R}}(\tilde{x}),\delta_{\tilde{M}}(\tilde{x})\delta_{\tilde{R}}(\tilde{x})\}\rangle=\text{L.H.S.}$ $\therefore, (\widetilde{\mathbf{L}} \vee_{\widetilde{Q}} \widetilde{M}) \boxplus \widetilde{R} = (\widetilde{\mathbf{L}} \boxplus \widetilde{R}) \vee_{\widetilde{Q}} (\widetilde{M} \boxplus \widetilde{R}).$ Hence, (ii) proved. $\mathrm{L.H.S} = \left(\tilde{\boldsymbol{L}} \wedge_{\widetilde{\boldsymbol{O}}} \widetilde{\boldsymbol{M}}\right) \boxtimes \tilde{\boldsymbol{R}}$ (iii) $= \langle \min\{\sigma_{\tilde{L}}(\tilde{x}), \sigma_{\tilde{M}}(\tilde{x})\}, \max\{\delta_{\tilde{L}}(\tilde{x}), \delta_{\tilde{M}}(\tilde{x})\} \rangle \boxplus \langle \sigma_{\tilde{R}}(\tilde{x}), \delta_{\tilde{R}}(\tilde{x}) \rangle$ $= \langle \min\{\sigma_{\tilde{L}}(\tilde{x}), \sigma_{\tilde{M}}(\tilde{x})\}\sigma_{\tilde{R}}, \left\{\max\left\{\delta_{\tilde{L}}(\tilde{x})^{\frac{5}{4}}, \delta_{\tilde{M}}(\tilde{x})^{\frac{5}{4}}\right\} + (\delta_{\tilde{R}}(\tilde{x}))^{\frac{5}{4}} - \max\left\{\delta_{\tilde{L}}(\tilde{x})^{\frac{5}{4}}, \delta_{\tilde{M}}(\tilde{x})^{\frac{5}{4}}\right\} (\delta_{\tilde{R}}(\tilde{x}))^{\frac{5}{4}}\right\}^{\frac{5}{5}} \rangle$ $= \langle \min\{\sigma_{\tilde{L}}(\tilde{x})\sigma_{\tilde{R}}(\tilde{x}),\sigma_{\tilde{M}}(\tilde{x})\sigma_{\tilde{R}}(\tilde{x})\}, \left\{\left(1-\left(\delta_{\tilde{R}}(\tilde{x})\right)^{\frac{5}{4}}\right)\max\left\{\delta_{\tilde{L}}(\tilde{x})^{\frac{5}{4}},\delta_{\tilde{M}}(\tilde{x})^{\frac{5}{4}}\right\}+\left(\delta_{\tilde{R}}(\tilde{x})\right)^{\frac{5}{4}}\right\}^{\frac{7}{5}}\rangle$ R.H.S = $(\tilde{L} \boxtimes \tilde{R}) \land_{\tilde{o}} (\tilde{M} \boxtimes \tilde{R})$ $= \langle \sigma_{\tilde{L}}(\tilde{x}) \sigma_{\tilde{R}}(\tilde{x}), \left\{ (\delta_{\tilde{L}}(\tilde{x}))^{\frac{5}{4}} + (\delta_{\tilde{R}}(\tilde{x}))^{\frac{5}{4}} - (\delta_{\tilde{L}}(\tilde{x}))^{\frac{5}{4}} (\delta_{\tilde{R}}(\tilde{x}))^{\frac{5}{4}} \right\}^{\frac{7}{5}} \rangle \wedge_{\tilde{0}}$ $\langle \sigma_{\tilde{M}}(\tilde{x})\sigma_{\tilde{R}}(\tilde{x}), \left\{ (\delta_{\tilde{M}}(\tilde{x}))^{\frac{5}{4}} + (\delta_{\tilde{R}}(\tilde{x}))^{\frac{5}{4}} - (\delta_{\tilde{M}}(\tilde{x}))^{\frac{5}{4}} (\delta_{\tilde{R}}(\tilde{x}))^{\frac{5}{4}} \right\}^{\frac{7}{5}} \rangle$ $= \langle min\{\sigma_{\tilde{L}}(\tilde{x})\sigma_{\tilde{R}}(\tilde{x}), \sigma_{\tilde{R}}(\tilde{x})\sigma_{\tilde{R}}(\tilde{x})\}, \{\left(1 - (\delta_{\tilde{R}}(\tilde{x}))^{\frac{5}{4}}\right) max\{\delta_{\tilde{L}}(\tilde{x})^{\frac{5}{4}}, \delta_{\tilde{M}}(\tilde{x})^{\frac{5}{4}}\} + (\delta_{\tilde{R}}(\tilde{x}))^{\frac{5}{4}}\}^{\frac{7}{5}}\} = \text{L.H.S.}$ $\therefore, (\tilde{\mathbf{L}} \wedge_{\tilde{\mathbf{O}}} \tilde{\mathbf{M}}) \boxtimes \tilde{\mathbf{R}} = (\tilde{\mathbf{L}} \boxtimes \tilde{\mathbf{R}}) \wedge_{\tilde{\mathbf{O}}} (\tilde{\mathbf{M}} \boxtimes \tilde{\mathbf{R}}).$ Hence, (iii) proved. Hence, $(\tilde{L} \wedge_{\tilde{Q}} \tilde{M}) \boxtimes \tilde{R} = (\tilde{L} \boxtimes \tilde{R}) \wedge_{\tilde{Q}} (\tilde{M} \boxtimes \tilde{R}).$ (iv) L.H.S = $(\tilde{L} \vee_{\widetilde{O}} \widetilde{M}) \boxtimes \widetilde{R}$ $= \langle \max\{\sigma_{\tilde{L}}(\tilde{x}), \sigma_{\tilde{M}}(\tilde{x})\}, \min\{\delta_{\tilde{L}}(\tilde{x}), \delta_{\tilde{M}}(\tilde{x})\} \rangle \boxplus \langle \sigma_{\tilde{R}}(\tilde{x}), \delta_{\tilde{R}}(\tilde{x}) \rangle$ $=\langle \max\{\sigma_{\tilde{L}}(\tilde{x}),\sigma_{\tilde{M}}(\tilde{x})\}\sigma_{\tilde{R}}(\tilde{x}), \left\{\min\left\{\delta_{\tilde{L}}(\tilde{x})^{\frac{5}{4}},\delta_{\tilde{M}}(\tilde{x})^{\frac{5}{4}}\right\} + (\delta_{\tilde{R}}(\tilde{x}))^{\frac{5}{4}} - \min\left\{\delta_{\tilde{L}}(\tilde{x})^{\frac{5}{4}},\delta_{\tilde{M}}(\tilde{x})^{\frac{5}{4}}\right\}(\delta_{\tilde{R}}(\tilde{x}))^{\frac{5}{4}}\right\}^{\frac{5}{5}}\rangle$ $= \langle \max\{\sigma_{\tilde{L}}(\tilde{x})\sigma_{\tilde{R}}(\tilde{x}),\sigma_{\tilde{M}}(\tilde{x})\sigma_{\tilde{R}}(\tilde{x})\}, \left\{\left(1-(\delta_{\tilde{R}}(\tilde{x}))^{\frac{5}{4}}\right)\min\left\{\delta_{\tilde{L}}(\tilde{x})^{\frac{5}{4}},\delta_{\tilde{M}}(\tilde{x})^{\frac{5}{4}}\right\}+(\delta_{\tilde{R}}(\tilde{x}))^{\frac{5}{4}}\right\}^{\frac{1}{5}}\rangle$ $R.H.S = (\tilde{\boldsymbol{L}} \boxtimes \tilde{R}) \vee_{\tilde{\boldsymbol{O}}} (\tilde{M} \boxtimes \tilde{R})$ $=\langle \sigma_{\tilde{L}}(\tilde{x})\sigma_{R}(\tilde{x}), \left\{ (\delta_{\tilde{L}}(\tilde{x}))^{\frac{5}{4}} + (\delta_{R}(\tilde{x}))^{\frac{5}{4}} - (\delta_{\tilde{L}}(\tilde{x}))^{\frac{5}{4}} (\delta_{R}(\tilde{x}))^{\frac{5}{4}} \right\}^{\frac{4}{5}} \vee \langle \sigma_{\tilde{M}}(\tilde{x})\sigma_{R}(\tilde{x}), \left\{ (\delta_{\tilde{M}}(\tilde{x}))^{\frac{5}{4}} + (\delta_{R}(\tilde{x}))^{\frac{5}{4}} - (\delta_{\tilde{M}}(\tilde{x}))^{\frac{5}{4}} (\delta_{R}(\tilde{x}))^{\frac{5}{4}} \right\}^{\frac{4}{5}} \rangle$ $= \langle \max\{\sigma_{\tilde{L}}(\tilde{x})\sigma_{\tilde{R}}(\tilde{x}),\sigma_{\tilde{M}}(\tilde{x})\sigma_{\tilde{R}}(\tilde{x})\},\min\left\{\left\{\left(1-(\delta_{\tilde{R}}(\tilde{x}))^{\frac{5}{4}}\right)(\delta_{\tilde{L}}(\tilde{x}))^{\frac{5}{4}}+(\delta_{\tilde{R}}(\tilde{x}))^{\frac{5}{4}}\right\}^{\frac{4}{5}},\left\{\left(1-(\delta_{\tilde{R}}(\tilde{x}))^{\frac{5}{4}}\right)(\delta_{\tilde{M}}(\tilde{x}))^{\frac{5}{4}}+(\delta_{\tilde{R}}(\tilde{x}))^{\frac{5}{4}}\right\}^{\frac{4}{5}}\right\}$ $=\langle \max\{\sigma_{\tilde{L}}(\tilde{x})\sigma_{R}(\tilde{x}),\sigma_{\tilde{M}}(\tilde{x})\sigma_{R}(\tilde{x})\}, \left\{\left(1-(\delta_{R}(\tilde{x}))^{\frac{5}{4}}\right)\min\left\{\delta_{\tilde{L}}(\tilde{x})^{\frac{5}{4}},\delta_{\tilde{M}}(\tilde{x})^{\frac{5}{4}}\right\}+(\delta_{R}(\tilde{x}))^{\frac{5}{4}}\right\}^{\frac{7}{5}}\rangle = \text{L.H.S.}$ $\therefore, (\tilde{\mathbf{L}} \vee_{\tilde{\mathbf{O}}} \tilde{\mathbf{M}}) \boxtimes \tilde{\mathbf{R}} = (\tilde{\mathbf{L}} \boxtimes \tilde{\mathbf{R}}) \vee_{\tilde{\mathbf{O}}} (\tilde{\mathbf{M}} \boxtimes \tilde{\mathbf{R}}).$ Hence, (iv) proved.

Theorem 3.2.4

Let $\widetilde{L} = \{ \langle \sigma_{\widetilde{L}}, \delta_{\widetilde{L}} \rangle : \forall \widetilde{x} \in X \}$ and $\widetilde{M} = \{ \langle \sigma_{\widetilde{M}}, \delta_{\widetilde{M}} \rangle : \forall \widetilde{x} \in X \}$ represents two Quin-Terranean fuzzy numbers, then,

(i)
$$(\tilde{L} \vee_{\tilde{Q}} \tilde{M}) \ominus (\tilde{L} \wedge_{\tilde{Q}} \tilde{M}) = \tilde{L} \ominus \tilde{M} \text{ if } \sigma_{\tilde{L}} \geq \sigma_{\tilde{M}}, \ \delta_{\tilde{L}} \leq \min \{\delta_{\tilde{M}}, \frac{\delta_{\tilde{M}} \varphi_1}{\varphi_2}\}.$$

Hence, $(\tilde{L} \vee_{\tilde{Q}} \tilde{M}) \boxtimes \tilde{R} = (\tilde{L} \boxtimes \tilde{R}) \vee_{\tilde{Q}} (\tilde{M} \boxtimes \tilde{R}).$

$$(\mathrm{ii}) \qquad \left(\tilde{L} \wedge_{\widetilde{Q}} \widetilde{M}\right) \ominus \left(\tilde{L} \vee_{\widetilde{Q}} \widetilde{M}\right) = \widetilde{M} \ominus \widetilde{L} \text{ if } \sigma_{\widetilde{L}} \geq \sigma_{\widetilde{M}}, \ \delta_{\widetilde{L}} \leq \min \left\{\delta_{\widetilde{M}}, \frac{\delta_{\widetilde{M}} \varphi_1}{\varphi_2}\right\}$$

$$(\mathrm{iii}) \qquad \left(\tilde{L} \wedge_{\tilde{Q}} \widetilde{M}\right) \oslash \left(\tilde{L} \vee_{\tilde{Q}} \widetilde{M}\right) = \tilde{L} \oslash \widetilde{M} \text{ if } \sigma_{\tilde{L}} \leq \min \left\{\sigma_{\tilde{M}}, \frac{\sigma_{\tilde{M}} \varphi_{1}}{\varphi_{2}}\right\}, \ \delta_{\tilde{L}} \geq \delta_{\tilde{M}}.$$





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$$\text{(iv)} \qquad \left(\tilde{L} \vee_{\widetilde{Q}} \widetilde{M}\right) \oslash \left(\tilde{L} \wedge_{\widetilde{Q}} \widetilde{M}\right) = \tilde{L} \oslash \widetilde{M} \text{ if } \sigma_{\tilde{L}} \leq \min \left\{\sigma_{\tilde{M}}, \frac{\sigma_{\widetilde{M}} \varphi_1}{\varphi_2}\right\}, \ \delta_{\tilde{L}} \geq \delta_{\tilde{M}}.$$

Proof:

Consider two Quin-Terranean fuzzy numbers $\tilde{L} = \langle \sigma_L, \delta_{\tilde{L}} \rangle$ and $\tilde{M} = \langle \sigma_{\tilde{M}}, \delta_{\tilde{M}} \rangle$.

(i) Since,
$$\sigma_{\tilde{L}} \geq \sigma_{\tilde{M}}$$
, $\delta_{\tilde{L}} \leq min\left\{\delta_{\tilde{M}}, \frac{\delta_{\tilde{M}} \varphi_1}{\varphi_2}\right\}$, then,

$$\text{L.H.S=}\left(\widetilde{\mathbf{L}}\vee_{\widetilde{Q}}\widetilde{M}\right)\ominus\left(\widetilde{\mathbf{L}}\wedge_{\widetilde{Q}}\widetilde{M}\right)$$

 $= \langle \max\{\sigma_{\tilde{L}}, \sigma_{\tilde{M}}\}, \min\{\delta_{\tilde{L}}, \delta_{\tilde{M}}\} \rangle \bigcirc \langle \min\{\sigma_{\tilde{L}}, \sigma_{\tilde{M}}\}, \max\{\delta_{\tilde{L}}, \delta_{\tilde{M}}\} \rangle$

$$= \langle \left\{ \frac{(\max\{\sigma_{L}, \sigma_{\bar{M}}\})^{4^{5}} - (\min\{\sigma_{L}, \sigma_{\bar{M}}\})^{4^{5}}}{1 - (\min\{\sigma_{\bar{L}}, \sigma_{\bar{M}}\})^{4^{5}}} \right\}^{\frac{1}{4^{5}}}, \frac{\min\{\delta_{L}, \delta_{\bar{M}}\}}{\max\{\delta_{\bar{L}}, \delta_{\bar{M}}\}} \rangle$$

$$= \langle \left\{ \frac{\max\{\sigma_{\bar{L}}^{4^{5}}, \sigma_{\bar{M}}^{4^{5}}\} - \min\{\sigma_{\bar{L}}^{4^{5}}, \sigma_{\bar{M}}^{4^{5}}\}}{1 - \min\{\sigma_{\bar{L}}^{4^{5}}, \sigma_{\bar{M}}^{4^{5}}\}} \right\}^{\frac{1}{4^{5}}}, \frac{\delta_{\bar{L}}}{\delta_{\bar{M}}} \rangle$$

$$(\sigma_{\bar{L}}^{4^{5}}, \sigma_{\bar{L}}^{4^{5}}, \sigma_{\bar{M}}^{4^{5}}, \sigma_{\bar{M}}^{4^{5}})$$

$$= \langle \left\{ \frac{\sigma_{\tilde{L}}^{4^{5}} - \sigma_{\tilde{M}}^{4^{5}}}{1 - \sigma_{\tilde{M}}^{4^{5}}} \right\}^{\frac{1}{4^{5}}}, \frac{\delta_{\tilde{L}}}{\delta_{\tilde{M}}} \rangle$$

$$= \widetilde{\boldsymbol{L}} \odot \widetilde{M} = R.H.S$$

$$\therefore, (\widetilde{L} \vee_{\widetilde{Q}} \widetilde{M}) \ominus (\widetilde{L} \wedge_{\widetilde{Q}} \widetilde{M}) = \widetilde{L} \ominus \widetilde{M}.$$

Hence, (i) proved.

(ii) Similarly, we can prove
$$(\tilde{L} \vee_{\tilde{Q}} \tilde{M}) \oslash (\tilde{L} \wedge_{\tilde{Q}} \tilde{M}) = \tilde{L} \oslash \tilde{M}$$
 if $\sigma_{\tilde{L}} \leq min \left\{ \sigma_{\tilde{M}}, \frac{\sigma_{\tilde{T}_2} \varphi_1}{\varphi_2} \right\}$, $\delta_{\tilde{L}} \geq \delta_{\tilde{M}}$.

(iii) Since,
$$\sigma_{\tilde{L}} \leq min\left\{\sigma_{\tilde{M}}, \frac{\sigma_{\tilde{M}}\varphi_1}{\alpha_2}\right\}$$
, $\delta_{\tilde{L}} \geq \delta_{\tilde{M}}$, then,

(iii) Since,
$$\sigma_{\tilde{L}} \leq min\left\{\sigma_{\tilde{M}}, \frac{\sigma_{\tilde{M}}\varphi_{1}}{\varphi_{2}}\right\}$$
, $\delta_{\tilde{L}} \geq \delta_{\tilde{M}}$, then,
L.H.S = $(\tilde{L} \wedge_{\tilde{Q}} \tilde{M}) \oslash (\tilde{L} \vee_{\tilde{Q}} \tilde{M}) = \langle \min \{\sigma_{\tilde{L}}, \sigma_{\tilde{M}}\}, \max\{\delta_{\tilde{L}}, \delta_{\tilde{M}}\} \rangle \oslash \langle \max\{\sigma_{\tilde{L}}, \sigma_{\tilde{M}}\}, \min\{\delta_{\tilde{L}}, \delta_{\tilde{M}}\} \rangle$

$$= \langle \frac{\min\{\sigma_{\bar{L}}, \sigma_{\bar{M}}\}}{\max\{\sigma_{\bar{L}}, \sigma_{\bar{M}}\}}, \left\{ \frac{(\max\{\delta_{\bar{L}}, \delta_{\bar{M}}\})^{\frac{5}{4}} - (\min\{\delta_{\bar{L}}, \delta_{\bar{M}}\})^{\frac{5}{4}}}{1 - (\min\{\delta_{\bar{L}}, \delta_{\bar{M}}\})^{\frac{5}{4}}} \right\}^{\frac{4}{5}} \rangle$$

$$= \langle \frac{\sigma_{\bar{L}}}{\sigma_{\bar{M}}}, \left\{ \frac{\max\left\{\delta_{\bar{L}}^{\frac{5}{4}}, \delta_{\bar{M}}^{\frac{5}{4}}\right\} - \min\left\{\delta_{\bar{L}}^{\frac{5}{4}}, \delta_{\bar{M}}^{\frac{5}{4}}\right\}}{1 - \min\left\{\delta_{\bar{L}}^{\frac{5}{4}}, \delta_{\bar{M}}^{\frac{5}{4}}\right\}} \right\}^{\frac{4}{5}} \rangle$$

$$=\langle \frac{\sigma_{\tilde{L}}}{\sigma_{\tilde{M}}}, \left\{ \frac{\delta_{\tilde{L}^{\frac{5}{4}}} - \delta_{\tilde{M}^{\frac{5}{4}}}}{1 - \delta_{\tilde{M}^{\frac{5}{4}}}} \right\}^{\frac{4}{5}} \rangle$$

$$= \tilde{L} \bigcirc \tilde{M} = R.H.S$$

$$\therefore, (\widetilde{L} \wedge_{\widetilde{O}} \widetilde{M}) \oslash (\widetilde{L} \vee_{\widetilde{O}} \widetilde{M}) = \widetilde{L} \oslash \widetilde{M}.$$

Hence, (iii) proved.

(iv) Similarly, we can prove,
$$(\tilde{L} \vee_{\tilde{Q}} \tilde{M}) \oslash (\tilde{L} \wedge_{\tilde{Q}} \tilde{M}) = \tilde{L} \oslash \tilde{M} \text{ if } \sigma_{\tilde{L}} \leq \min \left\{ \sigma_{\tilde{M}}, \frac{\sigma_{\tilde{M}} \varphi_1}{\varphi_2} \right\}, \ \delta_{\tilde{L}} \geq \delta_{\tilde{M}}.$$

Theorem 3.2.5

Let $\tilde{L} = \langle \sigma_{\tilde{L}}, \delta_{\tilde{L}} \rangle$ and $\tilde{M} = \langle \sigma_{\tilde{M}}, \delta_{\tilde{M}} \rangle$ represents two Quin-Terranean fuzzy numbers, then,

(i)
$$(\tilde{L} \wedge_{\tilde{O}} \tilde{M}) \vee_{\tilde{O}} \tilde{M} = \tilde{M}$$

(ii)
$$(\tilde{L} \vee_{\tilde{O}} \tilde{M}) \wedge_{\tilde{O}} \tilde{M} = \tilde{M}$$

(iii)
$$(\tilde{L} \ominus \tilde{M}) \boxplus \tilde{M} = \tilde{L} \text{ if } \sigma_{\tilde{L}} \ge \sigma_{\tilde{M}}, \ \delta_{\tilde{L}} \le \min \left\{ \delta_{\tilde{M}}, \frac{\delta_{\tilde{M}} \varphi_1}{\varphi_2} \right\}.$$

(v)
$$(\tilde{L} \oslash \tilde{M}) \boxtimes \tilde{M} = \tilde{L} \text{ if } \sigma_{\tilde{L}} \leq \min \left\{ \sigma_{\tilde{M}}, \frac{\sigma_{\tilde{M}} \varphi_1}{\varphi_2} \right\}, \ \delta_{\tilde{L}} \geq \delta_{\tilde{M}}.$$

Proof:

Consider two Quin-Terranean fuzzy numbers $\tilde{\mathbf{L}} = \langle \sigma_{\tilde{L}}, \delta_{\tilde{L}} \rangle$ and $\tilde{M} = \langle \sigma_{\tilde{M}}, \delta_{\tilde{M}} \rangle$.

(i) L.H.S =
$$(\tilde{L} \wedge_{\tilde{O}} \tilde{M}) \vee_{\tilde{O}} \tilde{M}$$

$$= \langle \min \{ \sigma_{\tilde{L}}, \sigma_{\tilde{M}} \}, \max \{ \delta_{\tilde{L}}, \delta_{\tilde{M}} \} \rangle \vee_{\tilde{O}} \langle \sigma_{\tilde{M}}, \delta_{\tilde{M}} \rangle$$

=
$$\langle \max\{\min\{\sigma_{\tilde{L}}, \sigma_{\tilde{M}}\}, \sigma_{\tilde{M}}\}, \min\{\max\{\delta_{\tilde{L}}, \delta_{\tilde{M}}\}, \delta_{\tilde{M}}\}\rangle$$

$$=\langle \sigma_{\tilde{M}}, \delta_{\tilde{M}} \rangle$$

$$=\widetilde{M}=\text{R.H.S.}$$





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 $:, (\widetilde{L} \wedge_{\widetilde{O}} \widetilde{M}) \vee_{\widetilde{O}} \widetilde{M} = \widetilde{M}.$

Hence, (i) proved.

(ii) L.H.S =
$$(\tilde{L} \vee_{\widetilde{O}} \widetilde{M}) \wedge_{\widetilde{O}} \widetilde{M}$$

$$= \langle \max \{ \sigma_{\tilde{L}}, \sigma_{\tilde{M}} \}, \min \{ \delta_{\tilde{L}}, \delta_{\tilde{M}} \} \rangle \cap \langle \sigma_{\tilde{M}}, \delta_{\tilde{M}} \rangle$$

$$= \langle \min\{\max\{\sigma_{\bar{L}}, \sigma_{\bar{M}}\}, \sigma_{\bar{M}}\}, \max\{\min\{\delta_{\bar{L}}, \delta_{\bar{M}}\}, \delta_{\bar{M}}\} \rangle$$

$$=\langle \sigma_{\widetilde{M}}, \delta_{\widetilde{M}} \rangle$$

$$=\widetilde{M}=\text{R.H.S.}$$

$$\therefore, (\widetilde{L} \vee_{\widetilde{O}} \widetilde{M}) \wedge \widetilde{M} = \widetilde{M}.$$

Hence, (ii) proved.

(iii) Since,
$$\sigma_{\tilde{L}} \geq \sigma_{\tilde{M}}$$
, $\delta_{\tilde{L}} \leq min \left\{ \delta_{\tilde{M}}, \frac{\delta_{\tilde{M}} \varphi_1}{\varphi_2} \right\}$, then

$$\text{L.H.S} = \left(\widetilde{\boldsymbol{L}} \bigcirc \widetilde{\boldsymbol{M}}\right) \boxplus \widetilde{\boldsymbol{M}} = \left\langle \left\{ \frac{\sigma_{\widetilde{\boldsymbol{L}}}^{45} - \sigma_{\widetilde{\boldsymbol{M}}}^{45}}{1 - \sigma_{\widetilde{\boldsymbol{M}}}^{45}} \right\}^{\frac{1}{45}}, \frac{\delta_{\underline{\boldsymbol{L}}}}{\delta_{\widetilde{\boldsymbol{M}}}} \right\rangle \boxplus \left\langle \sigma_{\widetilde{\boldsymbol{M}}}, \delta_{\widetilde{\boldsymbol{M}}} \right\rangle$$

$$= \langle \left\{ \left[\left(\frac{\sigma_{\tilde{L}}^{4^{5}} - \sigma_{\tilde{M}}^{4^{5}}}{1 - \sigma_{\tilde{M}}^{4^{5}}} \right)^{\frac{1}{4^{5}}} \right]^{4^{5}} + \sigma_{\tilde{M}}^{4^{5}} - \left[\left(\frac{\sigma_{\tilde{L}}^{4^{5}} - \sigma_{\tilde{M}}^{4^{5}}}{1 - \sigma_{\tilde{M}}^{4^{5}}} \right)^{\frac{1}{4^{5}}} \right]^{4^{5}} \sigma_{\tilde{M}}^{4^{5}} \right\}^{\frac{1}{4^{5}}}, \frac{\delta_{\tilde{L}} \delta_{\tilde{M}}}{\delta_{\tilde{M}}} \rangle$$

$$= \langle \left\{ \left(\frac{{\sigma_{\tilde{L}}}^{4^{5}} - {\sigma_{\tilde{M}}}^{4^{5}}}{1 - {\sigma_{\tilde{M}}}^{4^{5}}} \right) + {\sigma_{\tilde{M}}}^{4^{5}} - \left(\frac{{\sigma_{\tilde{L}}}^{4^{5}} - {\sigma_{\tilde{M}}}^{4^{5}}}{1 - {\sigma_{\tilde{M}}}^{4^{5}}} \right) {\sigma_{\tilde{M}}}^{4^{5}} \right\}^{\frac{1}{4^{5}}}, \delta_{\tilde{L}} \rangle$$

$$= \langle \left\{ \frac{\sigma_{\bar{L}}^{4^{5}} - \sigma_{\bar{M}}^{4^{5}} + \sigma_{\bar{M}}^{4^{5}} - \sigma_{\bar{M}}^{4^{5}} \sigma_{\bar{M}}^{4^{5}} - \sigma_{\bar{L}}^{4^{5}} \sigma_{\bar{M}}^{4^{5}} + \sigma_{\bar{M}}^{4^{5}} \sigma_{\bar{M}}^{4^{5}}}{1 - \sigma_{\bar{M}}^{4^{5}}} \right\}^{\frac{1}{4^{5}}}, \delta_{\bar{L}} \rangle$$

$$= \langle \left\{ \frac{\sigma_{\bar{L}}^{4^5} \left(1 - \sigma_{\bar{M}}^{4^5} \right)}{\left(1 - \sigma_{\bar{M}}^{4^5} \right)} \right\}^{\frac{1}{4^5}}, \delta_{\bar{L}} \rangle = \langle \left\{ \sigma_{\bar{L}}^{4^5} \right\}^{\frac{1}{4^5}}, \delta_{\bar{L}} \rangle = \langle \sigma_{\bar{L}}, \delta_{\bar{L}} \rangle$$

$$= \tilde{L} = R.H.S.$$

$$\therefore$$
, $(\tilde{L} \ominus \tilde{M}) \boxplus \tilde{M} = \tilde{L}$.

Hence, (iii) proved.

(iv) Since,
$$\sigma_{\tilde{L}} \leq min\left\{\sigma_{\tilde{M}}, \frac{\sigma_{\tilde{M}}\varphi_1}{\varphi_2}\right\}$$
, $\delta_{\tilde{L}} \geq \delta_{\tilde{M}}$, then,

$$\text{L.H.S} = \left(\tilde{\boldsymbol{L}} \oslash \widetilde{\boldsymbol{M}}\right) \boxtimes \widetilde{\boldsymbol{M}} = \langle \frac{\sigma_{\tilde{\boldsymbol{L}}}}{\sigma_{\tilde{\boldsymbol{M}}}}, \left\{ \frac{\delta_{\tilde{\boldsymbol{L}}}^{\frac{5}{4}} - \delta_{\tilde{\boldsymbol{M}}}^{\frac{5}{4}}}{1 - \delta_{\tilde{\boldsymbol{M}}}^{\frac{5}{4}}} \right\}^{\frac{4}{5}} \rangle \boxtimes \langle \sigma_{\tilde{\boldsymbol{M}}}, \delta_{\tilde{\boldsymbol{M}}} \rangle$$

$$\begin{split} = \langle \frac{\sigma_{L}\sigma_{\tilde{M}}}{\sigma_{\tilde{M}}}, \left\{ \left[\frac{\delta_{\tilde{L}^{\frac{5}{4}}}^{\frac{5}{4}} - \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}}}{1 - \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}}} \right]^{\frac{5}{4}}^{\frac{5}{4}} + \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} - \left\{ \left[\frac{\delta_{\tilde{L}^{\frac{5}{4}}} - \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}}}}{1 - \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}}} \right]^{\frac{5}{4}}^{\frac{5}{4}} \delta_{\tilde{M}^{\frac{5}{4}}} - \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} + \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} + \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} + \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} \right]^{\frac{5}{4}} \\ = \langle \sigma_{\tilde{L}}, \left[\frac{\delta_{\tilde{L}^{\frac{5}{4}}} - \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} + \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} - \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} + \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} + \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} \right]^{\frac{5}{4}} \\ = \langle \sigma_{\tilde{L}}, \left[\frac{\delta_{\tilde{L}^{\frac{5}{4}}} - \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} + \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} - \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} + \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} \right]^{\frac{5}{4}} \\ = \langle \sigma_{\tilde{L}}, \left[\frac{\delta_{\tilde{L}^{\frac{5}{4}}} - \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} + \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} - \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} + \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} \right]^{\frac{5}{4}} \\ = \langle \sigma_{\tilde{L}}, \left[\frac{\delta_{\tilde{L}^{\frac{5}{4}}} - \delta_{\tilde{L}^{\frac{5}{4}}}^{\frac{5}{4}} + \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} - \delta_{\tilde{L}^{\frac{5}{4}}}^{\frac{5}{4}} \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} + \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} \delta_{\tilde{M}^{\frac{5}{4}}}^{\frac{5}{4}} \right]^{\frac{5}{4}} \\ = \langle \sigma_{\tilde{L}}, \frac{\delta_{\tilde{L}^{\frac{5}{4}}} - \delta_{\tilde{L}^{\frac{5}{4}}}^{\frac{5}{4}} \delta_{\tilde{L}^{\frac{5}{4}}}^{\frac{5}{4}} \delta_{\tilde{L}^{\frac{5}{4}}}^{\frac{5}{4}} \delta_{\tilde{L}^{\frac{5}{4}}}^{\frac{5}{4}}} \right]^{\frac{5}{4}} \\ = \langle \sigma_{\tilde{L}}, \frac{\delta_{\tilde{L}^{\frac{5}{4}}} - \delta_{\tilde{L}^{\frac{5}{4}}}^{\frac{5}{4}} \delta_{\tilde{L}^{\frac{5}{4}}}^{\frac{5}{4}} \delta_{\tilde{L}^{\frac{5}{4}}}^{\frac{5}{4}} \delta_{\tilde{L}^{\frac{5}{4}}}^{\frac{5}{4}}} \right]^{\frac{5}{4}} \\ = \langle \sigma_{\tilde{L}}, \frac{\delta_{\tilde{L}^{\frac{5}{4}}} - \delta_{\tilde{L}^{\frac{5}{4}}}^{\frac{5}{4}} \delta_{\tilde{L}^{\frac{5}{4}}}^{\frac{5}{4}} \delta_{\tilde{L}^{\frac{5}{4}}}^{\frac{5}{4}$$

$$\begin{split} &= \langle \sigma_{\tilde{L}}, \left\{ \frac{\delta_{\tilde{L}^{\frac{5}{4}}} \left(1 - \delta_{\tilde{M}^{\frac{5}{4}}} \right)}{\left(1 - \delta_{\tilde{M}^{\frac{5}{4}}} \right)} \right\}^{\frac{4}{5}} \rangle = \langle \sigma_{\tilde{L}}, \left\{ \delta_{\tilde{L}^{\frac{5}{4}}} \right\}^{\frac{4}{5}} \rangle = \langle \sigma_{\tilde{L}}, \delta_{\tilde{L}} \rangle \\ &= \tilde{L} = \text{R.H.S.} \end{split}$$

$$\therefore, (\widetilde{L} \ominus \widetilde{M}) \boxplus \widetilde{M} = \widetilde{L}.$$

Hence, (iv) proved.

Theorem 3.2.6

Let
$$\tilde{\boldsymbol{L}} = \langle \sigma_{\tilde{\boldsymbol{L}}}, \delta_{\tilde{\boldsymbol{L}}} \rangle$$
, $\tilde{\boldsymbol{M}} = \langle \sigma_{\tilde{\boldsymbol{M}}}, \delta_{\tilde{\boldsymbol{M}}} \rangle$ and $\tilde{\boldsymbol{R}} = \langle \sigma_{\tilde{\boldsymbol{R}}}, \delta_{\tilde{\boldsymbol{R}}} \rangle$ represents three Quin-Terranean fuzzy numbers, then,
(i) $(\tilde{\boldsymbol{L}} \wedge_{\tilde{\boldsymbol{Q}}} \tilde{\boldsymbol{M}}) \ominus \tilde{\boldsymbol{R}} = (\tilde{\boldsymbol{L}} \ominus \tilde{\boldsymbol{R}}) \wedge_{\tilde{\boldsymbol{Q}}} (\tilde{\boldsymbol{M}} \ominus \tilde{\boldsymbol{R}})$,

$$\sigma_{\tilde{R}} \leq min\{\sigma_{\tilde{L}}, \sigma_{\tilde{M}}\}, max\{\delta_{\tilde{L}}, \delta_{\tilde{M}}\} \leq \left\{\delta_{\tilde{R}}, \frac{\delta_{\tilde{R}}\varphi_{1}}{\varphi_{3}}, \frac{\delta_{\tilde{T}_{3}}\varphi_{2}}{\varphi_{3}}\right\}, \frac{min\left\{\sigma_{\tilde{L}}^{4^{5}}, \sigma_{\tilde{M}}^{4^{5}}\right\} - \sigma_{\tilde{T}_{3}}^{4^{5}}}{\left(1 - \sigma_{\tilde{R}}^{4^{5}}\right)} + \frac{max\left\{\delta_{\tilde{L}}^{\frac{5}{4}}, \delta_{\tilde{M}}^{\frac{5}{4}}\right\}}{\delta_{\tilde{R}}^{\frac{5}{4}}} \leq 1;$$



if



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(ii)
$$(\tilde{L} \vee_{\tilde{Q}} \tilde{M}) \ominus \tilde{R} = (\tilde{L} \ominus \tilde{R}) \vee_{\tilde{Q}} (\tilde{M} \ominus \tilde{R}),$$

(iii) if
$$\sigma_{\tilde{R}} \leq min\{\sigma_{\tilde{L}}, \sigma_{\tilde{M}}\}, max\{\delta_{\tilde{L}}, \delta_{\tilde{M}}\} \leq \left\{\delta_{\tilde{R}}, \frac{\delta_{\tilde{R}}\varphi_1}{\varphi_3}, \frac{\delta_{\tilde{R}}\varphi_2}{\varphi_3}\right\}, \frac{max\left\{\sigma_{\tilde{L}}^{45}, \sigma_{\tilde{M}}^{45}\right\} - \sigma_{\tilde{T}_3}^{45}}{\left(1 - \sigma_{\tilde{R}}^{45}\right)} + \frac{min\left\{\delta_{\tilde{L}}^{\frac{5}{4}}, \delta_{\tilde{M}}^{\frac{5}{4}}\right\}}{\delta_{\tilde{L}}^{\frac{5}{4}}} \leq 1;$$

(iv)
$$(\tilde{\mathbf{L}} \wedge_{\widetilde{\mathbf{Q}}} \tilde{\mathbf{M}}) \oslash \tilde{R} = (\tilde{\mathbf{L}} \oslash \tilde{R}) \wedge_{\widetilde{\mathbf{Q}}} (\tilde{\mathbf{M}} \oslash \tilde{R})$$
 if $\delta_{\tilde{R}} \leq min\{\delta_{\tilde{\mathbf{L}}}, \delta_{\widetilde{\mathbf{M}}}\}, max\{\sigma_{\tilde{\mathbf{L}}}, \sigma_{\widetilde{\mathbf{M}}}\} \leq \left\{\sigma_{\tilde{R}}, \frac{\sigma_{\tilde{R}} \varphi_{1}}{\varphi_{3}}, \frac{\sigma_{\tilde{R}} \varphi_{2}}{\varphi_{3}}\right\}, \frac{max\left\{\delta_{\tilde{\mathbf{L}}}^{\tilde{4}}, \delta_{\tilde{\mathbf{M}}}^{\tilde{4}}\right\} - \delta_{T_{3}}^{\tilde{4}}}{\left(1 - \delta_{\tilde{R}}^{\tilde{4}}\right)} + \frac{min\left\{\sigma_{\tilde{\mathbf{L}}}^{45}, \sigma_{\tilde{\mathbf{M}}}^{45}\right\}}{\sigma_{3}^{45}} \leq 1;$

$$(v) \qquad \left(\tilde{\boldsymbol{L}} \vee_{\widetilde{\boldsymbol{Q}}} \widetilde{\boldsymbol{M}}\right) \oslash \widetilde{\boldsymbol{R}} = \left(\tilde{\boldsymbol{L}} \oslash \widetilde{\boldsymbol{R}}\right) \vee_{\widetilde{\boldsymbol{Q}}} \left(\widetilde{\boldsymbol{M}} \oslash \widetilde{\boldsymbol{R}}\right), \text{ if } \delta_{\widetilde{\boldsymbol{R}}} \leq min\{\delta_{\widetilde{\boldsymbol{L}}}, \delta_{\widetilde{\boldsymbol{M}}}\}, max\{\sigma_{\widetilde{\boldsymbol{L}}}, \sigma_{\widetilde{\boldsymbol{M}}}\} \leq \left\{\sigma_{\widetilde{\boldsymbol{R}}}, \frac{\sigma_{\widetilde{\boldsymbol{R}}} \varphi_{1}}{\varphi_{3}}, \frac{\sigma_{\widetilde{\boldsymbol{R}}} \varphi_{2}}{\varphi_{3}}\right\}, \frac{min\left\{\delta_{\widetilde{\boldsymbol{L}}}^{\widetilde{\boldsymbol{4}}}, \delta_{\widetilde{\boldsymbol{M}}}^{\widetilde{\boldsymbol{4}}}\right\} - \delta_{T_{3}}^{\widetilde{\boldsymbol{4}}}}{\left(1 - \delta_{\widetilde{\boldsymbol{R}}}^{\widetilde{\boldsymbol{4}}}\right)} + \frac{max\left\{\sigma_{\widetilde{\boldsymbol{L}}}^{45}, \sigma_{\widetilde{\boldsymbol{M}}}^{45}\right\}}{\sigma_{\widetilde{\boldsymbol{R}}}^{45}} \leq 1;$$

Proof:

Let $\tilde{\mathbf{L}} = \langle \sigma_{\tilde{\mathbf{L}}}, \delta_{\tilde{\mathbf{L}}} \rangle$, $\tilde{\mathbf{M}} = \langle \sigma_{\tilde{\mathbf{M}}}, \delta_{\tilde{\mathbf{M}}} \rangle$ and $\tilde{\mathbf{R}} = \langle \sigma_{\tilde{\mathbf{R}}}, \delta_{\tilde{\mathbf{R}}} \rangle$ be three Quin-Terranean fuzzy numbers.

(i) Since,
$$\sigma_{\tilde{R}} \leq min\{\sigma_{\tilde{L}}, \sigma_{\tilde{M}}\}, max\{\delta_{\tilde{L}}, \delta_{\tilde{M}}\} \leq \left\{\delta_{\tilde{R}}, \frac{\delta_{\tilde{R}}\varphi_1}{\varphi_3}, \frac{\delta_{\tilde{R}}\varphi_2}{\varphi_3}\right\}, \frac{min\left\{\sigma_{\tilde{L}}^{4^5}, \sigma_{\tilde{M}}^{4^5}\right\} - \sigma_{\tilde{\tau}_3}^{4^5}}{\left(1 - \sigma_{\tilde{R}}^{4^5}\right)} + \frac{max\left\{\delta_{\tilde{L}}^{\frac{5}{4}}, \delta_{\tilde{M}}^{\frac{5}{4}}\right\}}{\delta_{\tilde{c}}^{\frac{5}{4}}} \leq 1.$$

$$\mathrm{L.H.S} = \left(\tilde{\mathbf{L}} \wedge_{\widetilde{\mathbf{O}}} \widetilde{\mathbf{M}}\right) \ominus \widetilde{\mathbf{R}} = \left\langle \min\left\{\sigma_{\tilde{\mathbf{L}}}, \sigma_{\widetilde{\mathbf{M}}}\right\}, \max\left\{\delta_{\tilde{\mathbf{L}}}, \delta_{\widetilde{\mathbf{M}}}\right\}\right\rangle \ominus \left\langle\sigma_{\tilde{\mathbf{R}}}, \delta_{\tilde{\mathbf{R}}}\right\rangle$$

$$\begin{split} &= \langle \left\{ \frac{\min \left\{ \sigma_{\tilde{L}}^{4^{5}}, \sigma_{\tilde{M}}^{4^{5}} \right\} - \sigma_{\tilde{R}}^{4^{5}} \right\}^{4^{5}}}{\left(1 - \sigma_{R}^{4^{5}} \right)} \right\}^{4^{5}}, \frac{\max \left\{ \delta_{\tilde{L}}, \delta_{\tilde{M}} \right\}}{\delta_{\tilde{R}}} \rangle \\ &= \langle \min \left\{ \left(\frac{\sigma_{\tilde{L}}^{4^{5}} - \sigma_{R}^{4^{5}}}{1 - \sigma_{R}^{4^{5}}} \right)^{\frac{1}{4^{5}}}, \left(\frac{\sigma_{\tilde{M}}^{4^{5}} - \sigma_{R}^{4^{5}}}{1 - \sigma_{R}^{4^{5}}} \right)^{\frac{1}{4^{5}}} \right\}, \max \left\{ \frac{\delta_{\tilde{L}}}{\delta_{\tilde{R}}}, \frac{\delta_{\tilde{M}}}{\delta_{\tilde{R}}} \right\} \rangle. \end{split}$$

$$\left(\tilde{\boldsymbol{L}} \ominus \tilde{R}\right) \wedge_{\tilde{\boldsymbol{Q}}} \left(\tilde{\boldsymbol{M}} \ominus \tilde{R}\right) = \langle \left\{ \left(\frac{\sigma_{\tilde{\boldsymbol{L}}}^{4^{5}} - \sigma_{\tilde{\boldsymbol{R}}}^{4^{5}}}{1 - \sigma_{\tilde{\boldsymbol{R}}}^{4^{5}}}\right)^{\frac{1}{4^{5}}}\right\}, \frac{\delta_{\tilde{\boldsymbol{L}}}}{\delta_{\tilde{\boldsymbol{R}}}} \rangle \wedge_{\tilde{\boldsymbol{Q}}} \left\langle \left(\frac{\sigma_{\tilde{\boldsymbol{M}}}^{4^{5}} - \sigma_{\tilde{\boldsymbol{R}}}^{4^{5}}}{1 - \sigma_{\tilde{\boldsymbol{R}}}^{4^{5}}}\right)^{\frac{1}{4^{5}}}, \frac{\delta_{\tilde{\boldsymbol{M}}}}{\delta_{\tilde{\boldsymbol{R}}}} \rangle \right\rangle$$

$$=\langle \min\left\{\left(\frac{\sigma_{\tilde{L}}^{4^{5}}-\sigma_{\tilde{R}}^{4^{5}}}{1-\sigma_{\tilde{R}}^{4^{5}}}\right)^{\frac{1}{4^{5}}},\left(\frac{\sigma_{\tilde{M}}^{4^{5}}-\sigma_{\tilde{R}}^{4^{5}}}{1-\sigma_{\tilde{R}}^{4^{5}}}\right)^{\frac{1}{4^{5}}}\right\},\max\left\{\frac{\delta_{\tilde{L}}}{\delta_{\tilde{R}}},\frac{\delta_{\tilde{M}}}{\delta_{\tilde{R}}}\right\}\rangle$$

$$:, (\tilde{\mathbf{L}} \wedge_{\tilde{\mathbf{Q}}} \overset{\triangleright}{\mathbf{M}}) \ominus \tilde{\mathbf{R}} = (\tilde{\mathbf{L}} \ominus \tilde{\mathbf{R}}) \wedge_{\tilde{\mathbf{Q}}} (\tilde{\mathbf{M}} \ominus \tilde{\mathbf{R}})$$

Hence, (i) proved.

(ii) Similarly we can prove
$$(\tilde{L}V_{\tilde{Q}}\tilde{M}) \ominus \tilde{R} = (\tilde{L} \ominus \tilde{R})V_{\tilde{Q}}(\tilde{M} \ominus \tilde{R})$$
, if $\sigma_{\tilde{R}} \leq min\{\sigma_{\tilde{L}}, \sigma_{\tilde{M}}\}, max\{\delta_{\tilde{L}}, \delta_{\tilde{M}}\} \leq \{\delta_{\tilde{R}}, \frac{\delta_{\tilde{R}}\varphi_{1}}{\varphi_{3}}, \frac{\delta_{\tilde{R}}\varphi_{2}}{\varphi_{3}}\}, \frac{max\{\sigma_{\tilde{L}}^{45}, \sigma_{\tilde{M}}^{45}\} - \sigma_{\tilde{R}}^{45}}{(1 - \sigma_{\tilde{R}}^{45})} + \frac{min\{\delta_{\tilde{L}}^{\frac{5}{4}}, \delta_{\tilde{M}}^{\frac{5}{4}}\}}{\delta_{z}^{\frac{5}{4}}} \leq 1.$

(iii) Since,
$$\delta_{\tilde{R}} \leq min\{\delta_{\tilde{L}}, \delta_{\tilde{M}}\}, max\{\sigma_{\tilde{L}}, \sigma_{\tilde{M}}\} \leq \left\{\sigma_{\tilde{R}}, \frac{\sigma_{\tilde{R}}\varphi_{1}}{\varphi_{3}}, \frac{\sigma_{\tilde{R}}\varphi_{2}}{\varphi_{3}}\right\}, \frac{max\left\{\delta_{\tilde{L}}^{\frac{5}{4}}, \delta_{\tilde{M}}^{\frac{5}{4}}\right\} - \delta_{\tilde{R}}^{\frac{5}{4}}}{\left(1 - \delta_{\tilde{R}}^{\frac{5}{4}}\right)} + \frac{min\left\{\sigma_{\tilde{L}}^{45}, \sigma_{\tilde{M}}^{45}\right\}}{\sigma_{\tilde{R}}^{45}} \leq 1.$$

$$\mathrm{L.H.S} = \left(\tilde{L} \wedge_{\widetilde{Q}} \widetilde{M}\right) \oslash \widetilde{R} = \left\langle \min \left\{ \sigma_{\tilde{L}}, \sigma_{\widetilde{M}} \right\}, \max \left\{ \delta_{\tilde{L}}, \delta_{\widetilde{M}} \right\} \right\rangle \oslash \left\langle \sigma_{\tilde{R}}, \delta_{\tilde{R}} \right\rangle$$

$$=\langle\frac{\min\{\sigma_{\tilde{L}},\sigma_{\tilde{M}}\}}{\sigma_{\tilde{R}}},\left\{\frac{\max\left\{\delta_{\tilde{L}}^{\frac{5}{4}},\delta_{\tilde{M}}^{\frac{5}{4}}\right\}-\delta_{\tilde{R}}^{\frac{5}{4}}\right\}}{\left(1-\delta_{\tilde{R}}^{\frac{5}{4}}\right)}\right\}^{\frac{4}{5}}\rangle$$

$$= \langle \min\left\{\frac{\sigma_{\tilde{L}}}{\sigma_{\tilde{R}}}, \frac{\sigma_{\tilde{M}}}{\sigma_{\tilde{R}}}\right\}, \max\left\{\left(\frac{\delta_{\tilde{L}}^{\frac{5}{4}} - \delta_{\tilde{R}}^{\frac{5}{4}}}{\left(1 - \delta_{\tilde{R}}^{\frac{5}{4}}\right)}\right)^{\frac{4}{5}}, \left(\frac{\delta_{\tilde{M}}^{\frac{5}{4}} - \delta_{\tilde{R}}^{\frac{5}{4}}}{\left(1 - \delta_{\tilde{R}}^{\frac{5}{4}}\right)}\right)^{\frac{4}{5}}\right\}\rangle$$

R.H.S=
$$(\tilde{\mathbf{L}} \oslash \tilde{\mathbf{R}}) \land_{\tilde{\mathbf{O}}} (\tilde{\mathbf{M}} \oslash \tilde{\mathbf{R}})$$

$$=\langle \frac{\sigma_{\tilde{L}}}{\sigma_{\tilde{R}}}, \left(\frac{\delta_{\tilde{L}}^{\frac{5}{4}} - \delta_{\tilde{R}}^{\frac{5}{4}}}{\left(1 - \delta_{\tilde{R}}^{\frac{5}{4}}\right)}\right)^{\frac{4}{5}} \rangle \wedge_{\tilde{Q}} \langle \frac{\sigma_{\tilde{M}}}{\sigma_{\tilde{R}}}, \left(\frac{\delta_{\tilde{M}}^{\frac{5}{4}} - \delta_{\tilde{R}}^{\frac{5}{4}}}{\left(1 - \delta_{\tilde{R}}^{\frac{5}{4}}\right)}\right)^{\frac{4}{5}} \rangle$$





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$$\therefore, (\tilde{L} \wedge_{\tilde{O}} \tilde{M}) \oslash \tilde{R} = (\tilde{L} \oslash \tilde{R}) \wedge_{\tilde{O}} (\tilde{M} \oslash \tilde{R})$$

Hence, (iii) proved.

(iv) Similarly we can prove
$$(\tilde{L} \vee_{\tilde{Q}} \tilde{M}) \oslash \tilde{R} = (\tilde{L} \oslash \tilde{R}) \vee_{\tilde{Q}} (\tilde{M} \oslash \tilde{R})$$
, if $\delta_{\tilde{R}} \leq min\{\delta_{\tilde{L}}, \delta_{\tilde{M}}\}, max\{\sigma_{\tilde{L}}, \sigma_{\tilde{M}}\} \leq \{\sigma_{\tilde{R}}, \frac{\sigma_{\tilde{R}} \varphi_{1}}{\varphi_{3}}, \frac{min\{\delta_{\tilde{L}}^{\frac{5}{4}}, \delta_{\tilde{M}}^{\frac{5}{4}}\} - \delta_{\tilde{T}_{3}}^{\frac{5}{4}}}{(1 - \delta_{\tilde{R}}^{\frac{5}{4}})} + \frac{max\{\sigma_{\tilde{L}}^{4}, \sigma_{\tilde{M}}^{4}\}}{\sigma_{\tilde{R}}^{4}} \leq 1.$

Quin-Terranean Arithmetic mean operations over Quin-Terranean fuzzy numbers

In this section, Quin-Terranean Arithmetic mean operation over Quin-Terranean fuzzy numbers is defined.

Definition 3.3.1

Let $\tilde{\mathbf{L}} = \langle \sigma_{\tilde{\mathbf{L}}}, \delta_{\tilde{\mathbf{L}}} \rangle$, $\tilde{\mathbf{M}} = \langle \sigma_{\tilde{\mathbf{M}}}, \delta_{\tilde{\mathbf{M}}} \rangle$ represents two Quin-Terranean fuzzy numbers, then the average operator on $\tilde{\mathbf{L}}$ and $\tilde{\mathbf{M}}$ is characterized as

$$\tilde{L}@\tilde{\mathbf{M}} = \langle \frac{\sigma_{\tilde{L}}^{4^5} + \sigma_{\tilde{\mathbf{M}}}^{4^5}}{2}, \frac{\delta_{\tilde{L}^{\frac{5}{4}}} + \delta_{\tilde{\mathbf{M}}}^{\frac{5}{4}}}{2} \rangle$$

Properties of Quin-Terranean Arithmetic mean operations on Quin Terranean fuzzy numbers

The several properties of the max –min and min – max operations on Quin Terranean fuzzy numbers are given as follows,

Theorem 3.4.1

Let $\tilde{\mathbf{L}} = \langle \sigma_{\tilde{L}}, \delta_{\tilde{L}} \rangle$, $\tilde{\mathbf{M}} = \langle \sigma_{\tilde{M}}, \delta_{\tilde{M}} \rangle$ and $\tilde{R} = \langle \sigma_{\tilde{R}}, \delta_{\tilde{R}} \rangle$ represents three Quin-Terranean fuzzy numbers, then,

(i)
$$(\tilde{L} \wedge_{\tilde{O}} \tilde{M}) @ \tilde{R} = (\tilde{L} @ \tilde{R}) \wedge_{\tilde{O}} (\tilde{M} @ \tilde{R})$$

(ii)
$$(\tilde{\mathbf{L}} \vee_{\tilde{O}} \tilde{\mathbf{M}}) @ \tilde{R} = (\tilde{\mathbf{L}} @ \tilde{R}) \vee_{\tilde{O}} (\tilde{\mathbf{M}} @ \tilde{R})$$

Proof:

Let $\tilde{\mathbf{L}} = \langle \sigma_{\tilde{\mathbf{L}}}, \delta_{\tilde{\mathbf{L}}} \rangle$, $\tilde{\mathbf{M}} = \langle \sigma_{\tilde{\mathbf{M}}}, \delta_{\tilde{\mathbf{M}}} \rangle$ and $\tilde{\mathbf{R}} = \langle \sigma_{\tilde{\mathbf{R}}}, \delta_{\tilde{\mathbf{R}}} \rangle$ represents three Quin-Terranean fuzzy numbers, then,

(i) L. H. S =
$$(\tilde{L} \wedge_{\tilde{O}} \tilde{M}) @ \tilde{R}$$

=
$$\langle \min \{ \sigma_{\tilde{\tau}}, \sigma_{\tilde{m}} \}, \max \{ \delta_{\tilde{\tau}}, \delta_{\tilde{m}} \} \rangle @ \langle \sigma_{\tilde{\tau}}, \delta_{\tilde{\tau}} \rangle$$

$$= \langle \min \left\{ \frac{\sigma_{\tilde{L}}^{45} + \sigma_{\tilde{R}}^{45}}{2}, \frac{\sigma_{\tilde{M}}^{45} + \sigma_{\tilde{R}}^{45}}{2} \right\}, \max \left\{ \frac{\delta_{\tilde{L}}^{\frac{5}{4}} + \delta_{\tilde{R}}^{\frac{5}{4}}}{2}, \frac{\delta_{\tilde{M}}^{\frac{5}{4}} + \delta_{\tilde{R}}^{\frac{5}{4}}}{2} \right\} \rangle$$

$$=\langle \frac{{\sigma_{\tilde{L}}}^{4^{5}}+{\sigma_{\tilde{R}}}^{4^{5}}}{2}, \frac{{\delta_{\tilde{L}}}^{\frac{5}{4}}+{\delta_{\tilde{R}}}^{\frac{5}{4}}}{2} \rangle \wedge_{\tilde{Q}} \langle \frac{{\sigma_{\tilde{M}}}^{4^{5}}+{\sigma_{\tilde{R}}}^{4^{5}}}{2}, \frac{{\delta_{\tilde{M}}}^{\frac{5}{4}}+{\delta_{\tilde{R}}}^{\frac{5}{4}}}{2} \rangle$$

$$= (\tilde{L}@\tilde{R}) \wedge_{\tilde{O}} (\tilde{M}@\tilde{R}) = R. H. S$$

$$\therefore, (\widetilde{\mathbf{L}} \wedge_{\widetilde{O}} \widetilde{\mathbf{M}}) @ \widetilde{R} = (\widetilde{\mathbf{L}} @ \widetilde{R}) \wedge_{\widetilde{O}} (\widetilde{\mathbf{M}} @ \widetilde{R}).$$

Hence, (i) proved.

(ii) L. H. S =
$$(\tilde{L} \vee_{\tilde{O}} \tilde{M}) @ \tilde{R} = \langle \max \{ \sigma_{\tilde{L}}, \sigma_{\tilde{M}} \}, \min \{ \delta_{\tilde{L}}, \delta_{\tilde{M}} \} \rangle @ \langle \sigma_{\tilde{R}}, \delta_{\tilde{R}} \rangle$$

$$= \langle \max \left\{ \frac{\sigma_{\tilde{L}}^{4^{5}} + \sigma_{\tilde{R}}^{4^{5}}}{2}, \frac{\sigma_{\tilde{M}}^{4^{5}} + \sigma_{\tilde{R}}^{4^{5}}}{2} \right\}, \min \left\{ \frac{\delta_{\tilde{L}}^{\frac{5}{4}} + \delta_{\tilde{R}}^{\frac{5}{4}}}{2}, \frac{\delta_{\tilde{M}}^{\frac{5}{4}} + \delta_{\tilde{R}}^{\frac{5}{4}}}{2} \right\} \rangle$$

$$= \langle \frac{\sigma_{\tilde{L}}^{4^{5}} + \sigma_{\tilde{R}}^{4^{5}}}{2}, \frac{\delta_{\tilde{L}}^{\frac{5}{4}} + \delta_{\tilde{R}}^{\frac{5}{4}}}{2} \rangle \bigvee_{\tilde{Q}} \langle \frac{\sigma_{\tilde{M}}^{4^{5}} + \sigma_{\tilde{R}}^{4^{5}}}{2}, \frac{\delta_{\tilde{M}}^{\frac{5}{4}} + \delta_{\tilde{R}}^{\frac{5}{4}}}{2} \rangle \\ = (\tilde{L}@\tilde{R}) \bigvee_{\tilde{Q}} (\tilde{M}@\tilde{R}) = \text{R. H. S}$$

$$\therefore, \left(\tilde{\boldsymbol{L}} \vee_{\widetilde{\boldsymbol{Q}}} \tilde{\boldsymbol{M}}\right) @ \tilde{\boldsymbol{R}} = \left(\tilde{\boldsymbol{L}} @ \tilde{\boldsymbol{R}}\right) \vee_{\widetilde{\boldsymbol{Q}}} \left(\tilde{\boldsymbol{M}} @ \tilde{\boldsymbol{R}}\right).$$

Hence, (ii) proved.

Theorem 3.4.2

Let $\tilde{\mathbf{L}} = \langle \sigma_{\tilde{\mathbf{L}}}, \delta_{\tilde{\mathbf{L}}} \rangle$ and $\tilde{\mathbf{M}} = \langle \sigma_{\tilde{\mathbf{M}}}, \delta_{\tilde{\mathbf{M}}} \rangle$ represents two Quin-Terranean fuzzy numbers, then, $\left(\left(\tilde{\mathbf{L}} \wedge_{\tilde{Q}} \tilde{\mathbf{M}} \right) \boxplus \left(\tilde{\mathbf{L}} \vee_{\tilde{Q}} \tilde{\mathbf{M}} \right) \right) @ \left(\left(\tilde{\mathbf{L}} \wedge_{\tilde{Q}} \tilde{\mathbf{M}} \right) \boxtimes \left(\tilde{\mathbf{L}} \vee_{\tilde{Q}} \tilde{\mathbf{M}} \right) \right) = \tilde{\mathbf{L}}@\tilde{\mathbf{M}}.$





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Proof:

We have, min(a, b) + max(a, b) = a + b and $\min(a, b) \cdot \max(a, b) = a \cdot b$ Where a, b are two real numbers. Consider L.H.S, $(\tilde{L} \wedge_{\tilde{O}} \tilde{M}) \boxplus (\tilde{L} \vee_{\tilde{O}} \tilde{M}) = \langle \min\{\sigma_{\tilde{L}}, \sigma_{\tilde{M}}\}, \max\{\delta_{\tilde{L}}, \delta_{\tilde{M}}\} \rangle \boxplus \langle \max\{\sigma_{\tilde{L}}, \sigma_{\tilde{M}}\}, \min\{\delta_{\tilde{L}}, \delta_{\tilde{M}}\} \rangle$ $= \langle \left\{ \min\{\sigma_{\tilde{L}}^{4^5}, \sigma_{\tilde{\boldsymbol{M}}}^{4^5}\} + \max\{\sigma_{\tilde{L}}^{4^5}, \sigma_{\tilde{\boldsymbol{M}}}^{4^5}\} - \min\{\sigma_{\tilde{L}}^{4^5}, \sigma_{\tilde{\boldsymbol{M}}}^{4^5}\} \max\{\sigma_{\tilde{L}}^{4^5}, \sigma_{\tilde{\boldsymbol{M}}}^{4^5}\} \right\}^{\frac{1}{4^5}}, \max\{\delta_{\tilde{L}}, \delta_{\tilde{\boldsymbol{M}}}\} \min\{\delta_{\tilde{L}}, \delta_{\tilde{\boldsymbol{M}}}\} \rangle$ $= \langle \left\{ \sigma_{\tilde{L}}^{4^5} + \sigma_{\tilde{M}}^{4^5} - \sigma_{\tilde{L}}^{4^5} \sigma_{\tilde{M}}^{4^5} \right\}^{\frac{1}{4^5}}, \delta_{\tilde{L}} \delta_{\tilde{M}} \rangle$ $\left(\tilde{L} \wedge_{\tilde{O}} \tilde{M}\right) \boxtimes \left(\tilde{L} \vee_{\tilde{O}} \tilde{M}\right) = \left\langle \min\{\sigma_{\tilde{L}}, \sigma_{\tilde{M}}\}, \max\{\delta_{\tilde{L}}, \delta_{\tilde{M}}\} \right\rangle \boxtimes$ $\langle \max\{\sigma_{\tilde{I}}, \sigma_{\tilde{M}}\}, \min\{\delta_{\tilde{I}}, \delta_{\tilde{M}}\} \rangle$ $=\langle \min\{\sigma_{\tilde{\boldsymbol{L}}},\sigma_{\tilde{\boldsymbol{M}}}\}\max\{\delta_{\tilde{\boldsymbol{L}}},\delta_{\tilde{\boldsymbol{M}}}\}, \left\{\max\left\{\delta_{\tilde{\boldsymbol{L}}^{\frac{5}{4}}},\delta_{\tilde{\boldsymbol{M}}^{\frac{5}{4}}}\right\}+\min\left\{\delta_{\tilde{\boldsymbol{L}}^{\frac{5}{4}}},\delta_{\tilde{\boldsymbol{M}}^{\frac{5}{4}}}\right\}-\max\left\{\delta_{\tilde{\boldsymbol{L}}^{\frac{5}{4}}},\delta_{\tilde{\boldsymbol{M}}^{\frac{5}{4}}}\right\}\min\left\{\delta_{\tilde{\boldsymbol{L}}^{\frac{5}{4}}},\delta_{\tilde{\boldsymbol{M}}^{\frac{5}{4}}}\right\}\right\}^{\frac{1}{5}}\rangle$ $=\langle \sigma_{\tilde{L}}\sigma_{\tilde{M}}, \left\{ \delta_{\tilde{L}}^{\frac{5}{4}} + \delta_{\tilde{M}}^{\frac{5}{4}} - \delta_{\tilde{L}}^{\frac{5}{4}} \delta_{\tilde{M}}^{\frac{5}{4}} \right\}^{\frac{1}{5}} \rangle.$ Therefore, $L.H.S = \left(\left(\widetilde{L} \wedge_{\widetilde{O}} \widetilde{M} \right) \boxplus \left(\widetilde{L} \vee_{\widetilde{O}} \widetilde{M} \right) \right) @ \left(\left(\widetilde{L} \wedge_{\widetilde{O}} \widetilde{M} \right) \boxtimes \left(\widetilde{L} \vee_{\widetilde{O}} \widetilde{M} \right) \right)$ $= \langle \left\{ \sigma_{\tilde{L}}^{4^5} + \sigma_{\tilde{M}}^{4^5} - \sigma_{\tilde{L}}^{4^5} \sigma_{\tilde{M}}^{4^5} \right\}^{\frac{1}{4^5}}, \delta_{\tilde{L}} \delta_{\tilde{M}} \rangle @$ $\langle \sigma_{\tilde{L}} \sigma_{\tilde{M}}, \left\{ \delta_{\tilde{L}}^{\frac{5}{4}} + \delta_{\tilde{M}}^{\frac{5}{4}} - \delta_{\tilde{L}}^{\frac{5}{4}} \delta_{\tilde{M}}^{\frac{5}{4}} \right\}^{\overline{5}} \rangle$ $=\langle \frac{\sigma_{\tilde{L}}^{4^5} + \sigma_{\tilde{M}}^{4^5}}{2}, \frac{\delta_{\tilde{L}^{\frac{3}{4}}} + \delta_{\tilde{M}^{\frac{3}{4}}}}{2} \rangle = \tilde{L}@\tilde{M} = \text{R.H.S}$

CONCLUSION

In this paper, we have defined various operators namely, $\operatorname{Max} - \operatorname{Min}(V_{\overline{Q}})$, $\operatorname{Min} - \operatorname{Max}(\Lambda_{\overline{Q}})$ operators and Average operator (@) on Quin – Terranean Fuzzy Numbers. Several properties of these $\operatorname{Max} - \operatorname{Min}(V_{\overline{Q}})$ and $\operatorname{Min} - \operatorname{Max}(\Lambda_{\overline{Q}})$ operators as well as in combination with the basic operation of complement and model operators such as $\boxplus, \boxtimes, \ominus, \oslash$ already defined on Quin – Terranean Fuzzy Numbers are also studied. Additionally, several properties of the Average operator (@) in combination with $\operatorname{Max} - \operatorname{Min}(V_{\overline{Q}})$, $\operatorname{Min} - \operatorname{Max}(\Lambda_{\overline{Q}})$ operators and model operators on Quin-Terranean Fuzzy numbers are also analyzed. Quin-Terranean Fuzzy Numbers have more applications, and it would be possible to study their features. Furthermore, these operations on Quin-Terranean Fuzzy Numbers have applications in numerous other fields, such as decision-making, agriculture, and healthcare, etc which will be planned as a future work.

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RESEARCH ARTICLE

Artificial Intelligence's Place in Radiology: Current uses and Future **Prospects**

Naresh Kumar^{1*}, Bimal Nepal² and Dinesh Kumar³ and Manvee Rai⁴

- ¹Assistant Professor, Department of Radiology, School of Health Science, Om Sterling Global University, Hisar, Haryana, India.
- ²Research Scholar, Department of Radio Imaging Technology, College of Paramedical Sciences, Teerthanker Mahaveer University, Uttar Pradesh, India.
- ³Associate Professor, School of Pharmacy, Desh Bhagat University, Punjab, India.
- ⁴Assistant Professor, School of Paramedical Sciences ,Lifeline Educational Institute Sameda,Azamgarh U.P

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*Address for Correspondence

Naresh Kumar

Assistant Professor, Department of Radiology, School of Health Science, Om Sterling Global University, Hisar, Haryana, India. Email: nareshkumar59239@gmai.com



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ABSTRACT

This review explores how artificial intelligence (AI) has transformed radiology, redefining how medical images are analyzed and used. We delve into current AI applications, including image segmentation, lesion detection, disease classification, and image enhancement, all powered by advanced machine learning and deep learning techniques. Real-world case studies highlight AI's impact on diagnostic accuracy and operational efficiency. We also explain the core methodologies driving these radiological advancements, including deep learning architectures and convolutional neural networks. While AI holds promise, ethical concerns, biases in training data, and potential overreliance on automation necessitate careful consideration and collaboration between AI and human experts. Looking forward, we envision AI contributing to personalized treatment strategies through innovations like multimodal fusion and image-guided interventions. However, successful integration requires addressing challenges such as data quality and regulatory compliance, and ongoing training of radiologists to fully leverage Al's potential in clinical practice. We aim to illuminate the path that radiology has embarked upon - one where the collaboration between human expertise and AI-driven insights stands set to redefine the boundaries of radiologic imaging and patient care. In conclusion, this review underscores AI's transformative role in radiology, discussing current applications, methodologies, challenges, and prospects. This synergy





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between AI and human expertise has the capability to reshape medical imaging and enhance health care service.

Keywords: Artificial Intelligence (AI), Convolutional Neural Networks (CNNs), natural language processing (NLP), Natural Language Processing (NLP), picture archiving and communication systems (PACS).

INTRODUCTION

AI refers to the emulation of human intelligence processes by machines, particularly computer systems. Among these processes are learning, logical reasoning, problem-solving, perception, and language comprehension[1]. In recent years, Artificial intelligence (AI) is developing quickly, and this has caused a paradigm shift in the field of radiology [2]. As medical imaging technologies continue to produce increasingly intricate datasets, the incorporation of AI-driven methodologies has enabled a more nuanced and efficient interpretation of these images[3, 4]. Radiologists have long grappled with the intricate task of analyzing a wealth of medical images, extracting critical information to aid accurate diagnoses and treatment plans[5, 6, 7, 8, 9]. However, this task has grown in complexity as imaging modalities evolve and generate voluminous data that demands precise and swift analysis[10, 11]. The utilization of AI methods, especially machine learning and deep learning algorithms, has appeared as a ground-breaking solution. By harnessing the computational power of AI, radiologists are now empowered to decipher patterns, detect anomalies, and enhance the diagnostic process with unprecedented accuracy[12, 13, 14]. Beyond the realm of diagnostics, AI's impact resonates across various facets of radiology. From the early detection of diseases to the fine-tuning of treatment strategies, the synergy between AI and medical imaging has far-reaching implications for patient care[15, 16]. Nevertheless, this technological amalgamation comes with its set of difficulties.

Ethical considerations, data privacy concerns, and the potential for bias within AI algorithms underscore the need for a comprehensive exploration of the interplay between AI and radiology. A notable advancement involves the utilization of contemporary computer vision techniques, driven by deep learning, in the realm of medical applications. This progress is particularly concentrated on enhancing medical imaging, medical video analysis, and facilitating clinical deployment. One notable instance is the application of computer vision in the medical domain, as depicted in consider Fig 1, depicting a deep learning setup designed to extract valuable information simultaneously from two sources: images, typically analyzed using convolutional networks, and non-image data, typically managed by more versatile deep networks. The acquired explanation may encompass pathology diagnostics, symptoms, clinical forecasting, and various amalgamation. Fig 2 illustrates that CNNs can learn to generate images through training. Tasks vary, spanning from regression between images (as shown) to enhancing image resolution, creating new images, and various other objectives. Computer vision in clinical settings covers a wide range of tasks, including screening, diagnosis, condition detection, outcome prediction, and segmentation for identifying abnormalities within organs at the cellular level. It also involves monitoring diseases and providing support for clinical research. By generating accurate diagnostic results, they prevent human errors, ultimately saving lives and reducing expenses. Predictive and early diagnostics benefit from AI-enabled assistants that generate scan reports with high accuracy.AI is presently utilized in the research and development sphere, specifically in mining data from medical records. Cognitive technologies support healthcare organizations by utilizing extensive data resources to improve diagnostic capabilities[17].

The background and significance

The background and significance of the role of AI in radiology lies at the intersection of technological innovation and the evolving demands of modern healthcare. Historically, radiologists have relied on their expertise to analyse medical images and provide critical insights for diagnosis and treatment. But with the introduction of cutting-edge imaging technologies like computed tomography (CT), magnetic resonance imaging (MRI), and positron emission tomography (PET), data generation has increased dramatically, making it more challenging for radiologists to





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effectively process and interpret this data. This surge in data has created a bottleneck in the diagnostic process, leading to delays and the potential for missed diagnoses[18, 19,20]. In response, artificial intelligence (AI) technologies-in particular, machine learning and deep learning algorithms—have become a potent remedy. These algorithms have the capability to analyze extensive datasets and detect subtle patterns that might elude human observation. This ability has paved the way for breakthroughs in image segmentation, lesion detection, and disease classification. AI-driven image enhancement techniques further contribute to the extraction of clinically relevant information, facilitating more accurate and timely diagnoses[21]. The significance of this integration extends beyond diagnostic accuracy. Al's potential to streamline workflow processes, reduce interpretation times, and enhance efficiency holds the promise of addressing the challenges posed by the data deluge. Moreover, AI introduces the concept of predictive analytics, where algorithms can forecast disease progression and tre-atment responses, offering personalized and targeted interventions[22,23]. In essence, the integration of AI in radiology not only addresses the pressing challenges of data overload but also ushers in a new era of precision medicine. By augmenting radiologists' capabilities, AI has the capability to transform healthcare delivery, providing more accurate diagnoses, improved patient outcomes, and ultimately contributing to a more efficient and effective healthcare system. As such, understanding the background and significance of AI's role in radiology is crucial to navigating its benefits and hurdles in the swiftly changing environment of medical imaging[24 – 28].

Current Applications

In the present landscape of radiology, the integration of artificial intelligence (AI) has permeated various dimensions, propelling the field into new realms of accuracy and efficiency. AI's current applications encompass a spectrum of tasks, each addressing critical challenges within medical imaging.

Image Segmentation

AI algorithms excel at precisely delineating anatomical structures and identifying regions of interest within medical images. For instance, in brain MRI scans, AI-driven segmentation accurately separates different brain regions, aiding in the assessment of neurodegenerative disorders.

Lesion Detection

The ability of AI to detect subtle abnormalities is a game-changer. In mammography, AI-powered systems assist radiologists in spotting early signs of breast cancer by highlighting potential lesions that might otherwise be missed, thus enhancing diagnostic sensitivity.

Disease Classification

Al's capacity for pattern recognition has found prominence in disease classification. In the case of lung nodules detected in CT scans, AI algorithms can differentiate benign from malignant nodules with remarkable accuracy, expediting the diagnostic process and minimizing unnecessary interventions.

Image Enhancement

AI has the capacity to enhance image quality, particularly in cases of noisy or degraded images. This technology aids in reducing artifacts in MRI images, leading to improved visualization of soft tissue structures[29 – 33]. Furthermore, the integration of AI augments the radiologist's workflow. AI-powered tools can prioritize the order in which images are reviewed, reducing interpretation time. Moreover, AI-driven systems can flag cases that require immediate attention, ensuring that critical findings are swiftly brought to the radiologist's notice[21].

METHODOLOGIES AND TECHNIQUES

The incorporation of AI into radiology depends on a range of advanced methodologies and techniques, each designed to harness the intricacies of medical imaging data. Prominent among these are machine learning, deep





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learning, convolutional neural networks (CNNs), and natural language processing (NLP), each delivering distinct capabilities to enhance diagnostics across various imaging modalities.

Machine Learning

This foundational AI technique involves training algorithms to discern patterns within data and formulate forecasts or choices. In radiology, it can be used to develop predictive models for disease outcomes based on historical patient data[34, 35].

Deep Learning

it is a kind of machine learning, uses layers of neural networks to analyse complicated patterns. Deep learning excels at feature extraction, enabling radiologists to uncover intricate details in images that might elude traditional methods[36, 37].

Convolutional Neural Networks (CNNs)

A cornerstone of deep learning, CNNs are particularly well-suited for image analysis. Through the hierarchical analysis of visual characteristics, CNNs have shown impressive achievements in endeavors such as image categorization, identifying abnormalities, and dividing images into segments across various modalities including X-ray, MRI, CT, and PET scans [38, 39,40].

Natural Language Processing (NLP)

While NLP is more commonly associated with text analysis, its integration with radiology involves interpreting unstructured text data, such as radiology reports. NLP algorithms can extract relevant information from these reports, aiding in the correlation of clinical findings and image data[41, 42]. Collectively, these AI techniques and their application to diverse imaging modalities converge to transform the radiological landscape. By unlocking deeper insights and refining diagnostic accuracy, they underscore the potential of AI to augment radiologists' capabilities, ultimately translating to improved patient care.

CHALLENGES AND LIMITATIONS

Challenges:

- 1. **Workflow Disruption:** Incorporating new AI tools can disrupt established clinical workflows, potentially leading to initial inefficiencies as radiologists adapt to these changes [43, 44, 45].
- 2. **Integration with Existing Systems:**It takes meticulous coordination and technological know-how to ensure that AI is seamlessly integrated with current PACS and electronic health records (EHR) [44, 46].
- 3. **Data Compatibility:** AI models demand a large, diverse dataset for training. However, compatibility issues between AI algorithms and various imaging devices and formats can hinder effective model training [47, 48].
- 4. **Model Validation:** The dynamic nature of medical data necessitates continuous validation of AI models to ensure their accuracy and reliability over time.

Limitations

While the integration of AI in radiology presents ground-breaking possibilities, it is accompanied by an array of challenges and limitations that warrant careful consideration.

- 1. **Data Quality:** Al's efficacy relies heavily on high-quality, diverse, and well-curated data. However, medical imaging datasets can sometimes be incomplete, noisy, or biased, potentially compromising the performance and generalization of AI models.
- 2. **Interpretability:** Deep learning models, though highly effective, frequently serve as "black boxes, "It makes it difficult to grasp how they come to their conclusions. This lack of interpretability hinders radiologists from fully trusting and comprehending AI-generated results[49, 50].





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- 3. **Bias:** Pre-existing biases within training data can inadvertently perpetuate biases in AI outputs. If AI models are trained on imbalanced datasets, they might exhibit disparities in performance across demographic groups, leading to inequitable healthcare outcomes[50, 51].
- 4. **Regulatory Compliance:** Incorporating AI into medical practice is subject to stringent regulatory frameworks to ensure patient safety and data privacy. Compliance with these regulations, like the Health Insurance Portability and Accountability Act (HIPAA) in the United States, requires meticulous design, validation, and monitoring of AI algorithms[50, 52].
- 5. **Role in Replacing Radiologists:** A prevailing concern is the perception that AI might replace radiologists entirely. While AI can expedite routine tasks, it's essential to recognize that the nuanced judgment, clinical context, and empathy that radiologists provide cannot be replicated by machines[53, 54, 55].
- 6. **Collaboration between AI and Human Experts**: The optimal approach lies in harnessing AI as a collaborative tool rather than a replacement. AI can assist radiologists by automating time-consuming tasks, allowing them to concentrate on intricate analyses and decision-making.
- 7. **Skill Gap and Training:** Integrating AI into radiology necessitates training radiologists to understand AI concepts and outputs. Bridging this knowledge gap is essential to maximize AI's potential and foster effective collaboration between humans and machines.
- 8. **Ethical Dilemmas:** The ethical considerations of AI in radiology extend to issues such as patient consent, transparency, and accountability. Ensuring patients are informed about AI's role in their diagnoses and treatment plans is crucial for maintaining trust[50].

Navigating these challenges requires a multidisciplinary approach involving radiologists, data scientists, ethicists, and regulators. The goal is to harness AI's capabilities while mitigating risks, preserving human expertise, and ultimately providing safer, more accurate, and equitable patient care. Recognizing AI as a complementary tool, one that augments human insight, paves the way for a harmonious future where technology enhances rather than replaces radiological expertise.

OPPORTUNITIES

- 1. **Efficiency Gains:** AI can automate regular work like image pre-processing and preliminary analysis, enabling doctors in radiology department to concentrate on more complicated cases and interpretations[50, 56].
- 2. **Enhanced Diagnosis:** Al's pattern recognition capabilities can lead to earlier and more accurate disease diagnoses, ultimately improving patient outcomes[50, 57].
- 3. **Predictive Analytics:** AI can predict disease progression and treatment responses, aiding in personalized treatment planning[58, 59].
- 4. **Standardization:** AI can assist in standardizing the interpretation of images, potentially reducing variability in diagnoses across different radiologists.

STRATEGIES FOR EFFECTIVE INTEGRATION:

- 1. **Clinical Engagement:** Involving radiologists and other healthcare professionals early in the AI implementation process ensures that AI tools align with clinical needs and workflows[59].
- 2. **Interoperability:** Implementing AI solutions that are compatible with various imaging devices, PACS, and EHR systems ensures seamless integration[60, 61].
- 3. **Training and Education:** Radiologists need training to understand AI outputs, interpret results, and collaborate effectively with AI algorithms[62, 63].
- 4. **Model Validation and Monitoring:** Continuous validation and monitoring of AI models are critical to ensure consistent and accurate performance. Regular updates should address evolving clinical needs and changing data patterns.
- 5. **Ethical Considerations:** Addressing patient consent, data privacy, and the ethical implications of using AI-generated insights is crucial to building trust and maintaining ethical standards[64, 65].
- 6. **Gradual Implementation:** Phased deployment of AI solutions allows radiologists to gradually adapt to changes, minimizing disruption to clinical workflows[62].





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7. **Collaboration:** Collaborative efforts between radiologists, data scientists, engineers, and regulatory experts facilitate a comprehensive approach to AI integration[66, 67,68].

FUTURE DIRECTIONS

The field of radiology is on a dynamic trajectory with the evolution of artificial intelligence (AI), presenting a landscape filled with transformative possibilities. Several key trends emerge as we look towards the future multimodal imaging fusion, which harnesses AI to combine data from various imaging methods for a more comprehensive understanding of disease, explainable AI, which enhance transparency and trust by revealing how algorithms make decisions, federated learning, enabling AI training across institutions while preserving data privacy, AI-powered image guided interventions for real-time precision during procedures, precision medicine through AI's analysis of diverse patient data, automated report generation for efficient communication in health care, predictive analytics to forecast disease progression, and global accessibility via cloud-based AI platforms. Interdisciplinary collaboration, and addressing ethical and regulatory challenges will be essential as we embark on this transformative journey, promising a revolution in patient care and personalized medicine within the field of regulatory[69, 70,71,72].

CONCLUSION

In summary, this review paper underscores the pivotal role of artificial intelligence in radiology department. It addresses the challenges of complex radio imaging data and the need for innovative solutions, with AI integration enhancing diagnostic accuracy and workflow efficiency. AI's current applications in radiology, such as image segmentation and disease classification, are showcased with real-world examples. However, challenges like data quality, interpretability, and bias must be addressed. Understanding between radiologists and AI developers is crucial, leveraging human expertise with AI insights. The future of AI lies in radiology includes trends like multimodal imaging fusion and personalized treatments, promising precision medicine and improved patient care. Ultimately, AI has the future that lies in potential to revolutionize radiology, leading for diagnoses and streamlined workflow, benefiting the patient.

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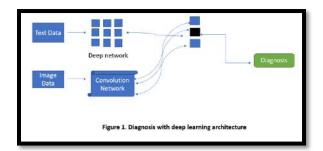


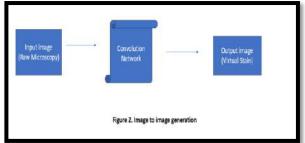
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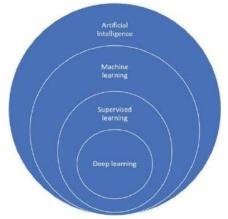


Figure 3: Components of Artificial Intelligence





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RESEARCH ARTICLE

The Prevalence of Anemia, along with its Associated Factors and **Treatment**

Umme Ayman^{1*}, Pallavi P¹, Sophy Thomas¹, Rahamtullah Khan¹, Manikanta BD¹ and Shailesh yadav²

¹Pharm.D. Intern, Department of Pharmacy Practice, Mallige College of Pharmacy, (Affiliated to Rajiv Gandhi University of Health Sciences) Bengaluru, Karnataka, India.

²HOD, Associate Professor, Department of Pharmacy Practice, Mallige College of Pharmacy, (Affiliated to Rajiv Gandhi University of Health Sciences) Bengaluru, Karnataka, India.

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*Address for Correspondence

Umme Ayman

Pharm.D. Intern, Department of Pharmacy Practice, Mallige College of Pharmacy, (Affiliated to Rajiv Gandhi University of Health Sciences) Bengaluru, Karnataka, India. Email: ainuainu75@gmail.com



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ABSTRACT

Anemia constitutes a formidable public health challenge that frequently goes undetected or untreated. Despite advancements in healthcare, it still afflicts a quarter of the global populace. This study aims to demonstrate the prevalence, severity, types, available treatment alternatives, and correlated factors of anemia among hospitalized inpatients at a tertiary care hospital in Bengaluru, Karnataka. We undertook a cross-sectional observational study design at a tertiary care hospital. To gather data, we looked at the medical records of the clinical, biochemical, and medication profiles of recently registered patients in the hospital from April 2022 to July 2022. The chi-square and p-value were employed to identify the association between anemia and associated factors. The study found that 15.31% of the population had anemia, with 35.9% mild, 47.1% moderate, and 17.1% severe anemia. Of the total anemic cases, 52.4% were males, while 47.6% were females, and more than half of the patients had normocytic anemia, i.e., normocytic 62.5%, microcytic 31.3%, and macrocytic 6.3%. As individuals grow older, the probability of developing anemia rises. Therefore, it is advisable to incorporate regular anemia screening into routine check-ups, particularly for high-risk groups who may have additional health concerns. Early recognition and intervention by healthcare providers can also lessen the number of untreated and under diagnosed anemic patients. Educating the public about iron- and vitamin-rich (vitamin B12 and B9) sources of plantand animal-based foods is an efficient strategy for reducing anemia prevalence in both vegetarian and non-vegetarian communities.





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Keywords: Anemia, gender, hemoglobin, prevalence, treatment.

INTRODUCTION

Anemia implies a decrease in erythrocyte production and quality, substantially impairing the body's ability to provide O2 to its tissues1. It is a symptom rather than a diagnosis, and its cause, severity, and multiple disorders, such as cardiovascular illness, influence symptoms. Once the Hb drops below 7.0 g/dL, symptoms like tiredness, weakness, and lethargy are common. Men, women, and children commonly lie within the range of 13.5-18.0 g/dL, 12-15 g/dL, and 11-16 g/dL, respectively. Pregnancy values vary depending on the trimester but are often larger than 10.0 g/dL². The corrected reticulocyte count categorizes anemia into two main types: hypoproliferative (below2%) and hyperproliferative (above2%). Microcytic (MCV is above 80 fl), normocytic (MCV is betwixt80 and 100 fl), and macrocytic (MCV is below 100 fl) anemias are additional classifications based on the mean corpuscular volume². The WHO has identified that roughly two billion populaces worldwide suffer from anemia, with iron deficiency being the leading cause in 50% of cases. Anemia is less common than iron deficiency by a factor of 2.5, with developing countries experiencing higher rates of anemia. Children under 5 have an estimated anemia prevalence of 39%, while the figures stand at 48% for children aged 5-14, 42% for women aged 15-59, 30% for men aged 15-59, and 45% for adults over 60 years. These statistics have significant health and economic implications for low- and middle-income countries[4]. Anemia can be upshot from reduced erythrocyte synthesis oran increased loss of blood via hemolysis, bleeding, or a combination of both. Nutritional, viral, or genetic factors may affect these. Hemoglobinopathies like sickle cell and thalassemia result from hereditary factors, while infectious disorders like malaria and schistosomiasis are common causes of anemia. Nutritional anemia results from a lack of minerals needed for hemoglobin production and erythropoiesis, including iron, vitamin B9, cobalamin, retinol, and protein-energy malnutrition. Iron deficiency is responsible for half of all anemias. Exposure to minor levels of trace elements, such as Zn and Cu, and hazardous heavy metals, such as lead, can potentially aggravate anemia[5]. Anemia is treatable by drug-based (iron tablets, folic acid, B12, and blood transfusions) and non-drug-based (vegetables, fruits, rosella beverages, and Moringa leaf juice) methods[6]. The main area of concern is the underlying etiology of anemia. For acute blood loss, blood transfusions, oxygen, and IV fluids are used, with a target of 7 g/dL for most patients. Oral or IV iron, B12, and folate all address nutritional deficiencies. Oral iron supplements are the most popular way to replenish iron, with the amount based on age, estimated iron deficiency, needed rate of correction, and tolerance to adverse effects. Bone marrow transplantation is required for a plastic anemia, while erythropoietin is effective for chronic diseases. Treatment of underlying diseases is necessary for autoimmune and rheumatological disorders. Increased red blood cell destruction results in hemolytic anemia, which calls for splenectomy, medicine, and mechanical valves. Hemoglobinopathies, namely sickle cell disease, necessitate blood transfusions, red cell exchange transfusions, and hydroxyurea. Bleeding that might be deadly is treated with antifibrinolytic medications[2].

MATERIALS AND METHODS

We undertook across-sectional observational study designat Mallige Medical Center, a multi-specialty tertiary care hospital in Bengaluru. The data available originated from the medical records of the clinical, biochemical, and pharmaceutical profiles of new patients admitted to the hospital between January 2022 and April 2022. The BMI was determined for each patient via available parameters. The inclusion and exclusion criteria are as follows: The study comprised all newly registered patients attending the hospital's inpatient department who were at least 18 years of age and older and whose records included a clinical history, complete blood count calculation, and medication chart during the study period. Patients who had just undergone surgery, pregnant women, and nursing mothers served as exclusion criteria. Before collecting any data, there view board of Mallige College of Pharmacy granted ethical approval and provided a letter of concurrence to Mallige Medical Center. The patient or a bystander was fully informed of the aim and purpose of the study, as required. A precise sample size of 1,110 patients who met established criteria was determined utilizing the single population proportion formula. To analyze the data, we





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employed Microsoft Excel for statistical tabulation. We ensured that the data was accurate and free of any missing values. We then manually reviewed the results with tallies and a scientific calculator. We performed a rigorous statistical analysis using the chi-square test with a significance level of 0.05 to determine the correlation between the independent and dependent variables.

RESULTS

Out of 1,110 respondents analyzed based on inclusion and exclusion criteria, 170 were acknowledged as anemic, resulting in an overall prevalence rate of 15.31%. Out of 170 patients, the percentages of mild, moderate, and severe anemia were 35.88%, 47.06%, and 17.06%, respectively. The distribution of various types of anemia among the study population showed 31.18% with microcytic anemia, 62.35% with normocytic anemia, and 6.48% with macrocytic anemia. Table 4 shows that 170 out of 1,110 patients had anemia, of whom 52.35% were male patients (n = 89) and 47.65% were female patients (n = 81). Among all age groups, young adult patients (18-30 years) had the lowest prevalence of anemia (4.12%). Furthermore, quadragenarians (41-50 years, 12.35%) and the elderly (81 years and older, 12.35%) had the lowest rates of anemia. According to our findings, the age category of 61-70 years had the highest proportion of anemia (22.94%). Anemia was higher among urban residents (81.20%) than rural residents (18.80%). Most patients had a normal BMI (57.65%), while 17.06% were underweight, 16.47% were obese, and 8.82% were overweight. About 89 (52.35%) of the anemic patients were vegetarians, whereas the other 81 (47.65%) were non-vegetarians. The prevalence of blood type A (31.18%) surpasses that of blood type B (30%), blood type AB (19.41%), and blood type O (19.41%) among anemic subjects. Age and anemia were significantly correlated (x^2 = 63.476, p = 0.00001). There was an association between BMI and anemia ($x^2 = 11.34$, p = 0.01002). It has been observed that anemia had a significant correlation with the blood group ($x^2 = 9.57$, p = 0.0226), but not statistically. Anemia had no significant association with gender ($x^2 = 1.215$, p = 0.2704), residence ($x^2 = 0.711$, p = 0.3991),or diet ($x^2 = 0.896$, p = 0.3438). Hypertension was discovered to be the most common co-morbidity in 170 anemic patients, accounting for 61.17%, followed by diabetes mellitus (41.76%), and others (i.e., 20%), including seizures, Crohn's disease, tuberculosis, obstructive sleep apnea, osteoporosis, and CAD (18.24%). Most patients who were mildly anemic received vitamin B supplements; just 3.28% received blood transfusions, and more than half (57.37%) did not get any treatment. Vitamin B supplements were administered to 37.5% of patients with moderate anemia, while blood transfusions were given to 20%, erythropoietin to 5%, vitamin B and iron supplements to 5%, and only iron supplements to 2.5%. Around 30% of moderately anemic patients went untreated. About 86.2% of severely anemic patients received blood transfusions, with 6.9% receiving vitamin B and iron supplements, 3.4% receiving iron supplements, and the remaining 3.4% receiving vitamin B supplements.

DISCUSSION

The study found a 15.31% overall prevalence of anemia. The observed prevalence is greater than the anemic prevalence of 15%reported at Hawas University in southern Ethiopia[7]. In contrast to the study done at the Gilgel Gibe Field Research Center in Ethiopia (40.9%), the prevalence found in this study is lower. This difference could be due to sample size, geographical variation, or socio-demographic variation[8]. Contrary to research done at Hawas University in southern Ethiopia, where a large percentage of individuals had mild anemia (58.5%), the majority of cases in this study had moderate anemia (47.06%)7. 17.06% of the participants had severe anemia, a rate greater than Yaounde-Cameroon's detection rate (0.8%)3 but lower than Hawas University's research in southern Ethiopia, which found 22.5% of participants to have severe anemia[7]. Age-related categories of anemia patients revealed that those over 51 had the highest prevalence (75.88%) of cases. Anemia and age were statistically linked, with severe cases being more prevalent in those aged 61 and above, who may also be affected by other health issues like hypertension, diabetes mellitus, and chronic kidney disease, all of which can impact the production of red blood cells. Anemia is more likely to occur as people get older because of various health conditions that may also be present. Males were slightly more common than females among the 170 anemic patients (52.35% vs. 20.3%), whereas research done in





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New Jersey, USA, found the opposite to be true (79.6% vs. 20.3%) than males(20.3%)7. Females are more likely to acquire anemia because of menstruation and childbirth. Gender was not significantly associated with anemia, which might be because, as age increases, both sexes are more likely to have comorbid conditions that increase the risk of anemia. Urban residents had a greater rate of anemia (81.20%) than rural residents (18.80%). The Gilgel Gibe Field Research Center in Ethiopia conducted a similar study that found that rural individuals were more susceptible to anemia than urban ones8. Anemia and residence had no significant association. The discrepancy in our results could be because our study was conducted solely in an urban setting, which makes it difficult to determine the exact distribution of anemia in urban and rural areas. The results of the study show a notable correlation between BMI and anemia. The majority of patients had a normal BMI (57.65%), followed by underweight (17.06%) and obese (16.47%) patients. Malnutrition is a common cause of low blood counts in underweight patients¹⁰, while obesity can interfere with iron hemostasis, leading to anemia¹¹. The study unequivocally demonstrates no significant link between diet and anemia. It is interesting to note that of the participants, 52.35% followed a vegetarian diet, while 47.65% were nonvegetarians. Vegetarians forego meat and eggs, which can result in iron and vitamin B12 deficiencies and, eventually, anemia¹³.Among 170 patients with anemia, hypertension was the most common complication, affecting 61.17% of patients. Diabetes mellitus was also prevalent in 41.76% of patients, while CAD was prevalent in 18.24%. A similar study conducted in Camden, New Jersey, USA, found that the most common complications were hypertension (59.3%), hypothyroidism (29.3%), and diabetes mellitus (19.4%)¹² There was a note worthy association between blood type and anemia. Among the 170 anemic patients studied, individuals with blood types A(31.18%) and B (30%) were observed to possess a higher susceptibility to anemia in contrast to those with blood types AB and O. Additionally, a similar study performed in North India revealed that the prevalence of anemia was maximum in blood type O(39.24%). Following this, blood type Baccounts for 42.20%, then blood type A for 13.3%, and blood type AB for 5.25%14. More than 50% of patients with mild anemia and 30% of those with moderate anemia were not appropriately diagnosed or treated. Although mild anemia is not a serious concern, if left untreated, it can cause serious health problems like tiredness and cardiac issues like arrhythmias, which may present a high risk of complications and, in rare cases, even death¹⁵.

CONCLUSION

As individuals grow older, the probability of developing anemia rises. Therefore, it is advisable to incorporate regular anemia screening into routine check-ups, particularly for high-risk groups who may have additional health concerns. Early recognition and intervention by healthcare providers can also lessen the number of untreated and under diagnosed anemic patients. Educating the public about iron- and vitamin-rich (vitamin B12 and B9) sources of plant- and animal-based foods is an efficient strategy for reducing anemia prevalence in both vegetarians and non-vegetarians.

ABBREVIATIONS

BMI: body mass index;

CAD: coronary artery disease; CHF: congestive heart failure; CKD: chronic kidney disease;

COPD: chronic obstructive pulmonary disease;

CVA: cerebrovascular accident;

Hb: hemoglobin;

MCV: mean corpuscular volume; WHO: World Health Organization





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Potential Conflict of Interest

There are no conflicting interests.

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Table 1. The WHO recommends hemoglobin values (g/dL) to determine anemia's severity in various populations³.

1 1				
Subject group	Normal	Mildly	Moderately	Severely
Subject group	Normal	anemic	anemic	anemic
Infants aged 6–59 months	≥ 11	10-10.9	7–9.9	<7
Children aged 5–11 years	≥ 11.5	11-11.4	8-10.9	<8
Adolescents aged 12–14 years	≥ 12	11-11.9	8-10.9	<8
Non-pregnant females aged 15 years and	≥ 12	11–11.9	8–10.9	<8
above	≥ 12	11-11.9	6-10.9	%
Gestational females	≥ 11	10-10.9	7–9.9	<7
Males aged 15 years and above	≥ 13	11–12.9	8-10.9	<8

Table 2. Prevalence and severity of anemia

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Grading of anemia severity	No. of Cases	Percentage (%)			
Mild	61	35.88			
Moderate	80	47.06			
Severe	29	17.06			
Total	170	100			

Table 3. Grading of anemia types among patients

Anemic types	No. of Cases	Percentage (%)
Microcytic	53	31.18
Normocytic	106	62.35
Macrocytic	11	6.47
Total	170	100

Table 4. Anemia-related factors among inpatients at a tertiary care hospital in Bengaluru (n=170)

Characteristics	Anemic subjects (n=170)	Percentage (%)	Chi-square(x²)	P-value				
	Gender							
Male	89	52.35	1.015					
Female	Female 81		1.215	0.2704				
Age category								
18–30years	7	4.12						
31–40years	13	7.65						
41–50years	21	12.35	63.476	0.00001				
51–60years	38	22.35						
61–70years	39	22.94						





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71–80years	31	18.24		
81 years and older	21	12.35		
	Residence			
Urban	138	81.20		
Rural	32	18.80	0.711	0.3991
	BMI category			
Underweight (less than 18.5)	29	17.06		
Healthy weight (18.5 to 24.9)	98	57.65		
Overweight (25 to 29.9)	15	8.82	11.34	0.01002
Obese (more than or equal to 30)	28	16.47		
	Diet			
Vegetarian	89	52.35	0.896	
Non-Vegetarian	81	47.65		0.3438
	Blood types			
A	53	31.18		
В	51	30	0.57	0.000
AB	33	19.41	9.57	0.0226
0	33	19.41		

Table 5. Prevalence of anemia based on co-morbidities

Co-morbidities	Anemic subjects (n =170)	Percentage (%)
Hypertension	104	61.17
Diabetes mellitus	71	41.76
COPD/Asthma	27	15.88
Thyroid illness	23	13.53
CKD	13	7.64
Malignancy	1	0.6
Rheumatological disease	1	0.6
CHF	1	0.6
CAD	31	18.24
CVA	9	5.3
Others	34	20
None	26	15.3

Table 6. Type of anemia and different options of treatment, showing the percentage distribution

Anemia Status	Vitamin B Supplements,n(%)	Iron Supplements,n(%)	Vitamin B+ Iron,n(%)	Blood Transfusion,n(%)	Erythropoietin,n(%)	None,n(%)	Total, n(%)
Mild Anemia							
Microcyti c	5 (8.19)	_	_	-	-	8 (13.11)	13 (21.3)





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Normocyt ic	18 (29.5)	-	1 (1.63)	2 (3.28)	-	26 (42.62)	47 (77)
Macrocyti c	-	-	-	-	-	1 (1.63)	1 (1.63)
Total	23 (37.7)	ı	1 (1.63)	2 (3.28)	-	35 (57.37)	61 (100)
		Mod	derate Anei	mia			
Microcyti c	7 (8.75)	1 (1.25)	3 (3.75)	6 (7.5)	2 (2.5)	6 (7.5)	25 (31.2 5)
Normocyt ic	19 (23.75)	1 (1.25)	1 (1.25)	10 (12.5)	2 (2.5)	17 (21.25)	50 (62.5)
Macrocyti c	4 (5)	-	-	-	-	1 (1.25)	5 (6.2)
Total	30 (37.5)	2 (2.5)	4 (5)	16 (20)	4 (5)	24 (30)	80 (100)
	Severe Anemia						
Microcyti c	-	1 (3.4)	-	11 (38)	-	_	12 (41.3 7)
Normocyt ic	1 (3.4)	-	-	8 (27.58)	-	_	9 (31.0 3)
Macrocyti c	-	-	2 (6.9)	6 (20.68)	_	_	8 (27.6)
Total	1 (3.4)	1 (3.4)	2 (6.9)	25 (86.2)	_	_	29 (100)

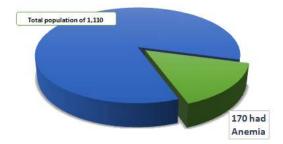


Figure 1.Prevalence of anemia





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REVIEW ARTICLE

Orodispersible Tablets: A Review

G.S.Sharma^{1*}, Namami Dutt², R. Shireesh Kiran and T. Rama Rao

¹Associate Professor, Department of Pharmaceutics, CMR College of Pharmacy, (Affiliated to Jawaharlal Nehru Technological University, Hyderabad) Telangana, India.

²Student, Department of Pharmaceutics, CMR College of Pharmacy, (Affiliated to Jawaharlal Nehru Technological University, Hyderabad) Telangana, India.

³Associate Professor, Department of Pharmaceutics, CMR College of Pharmacy, (Affiliated to Jawaharlal Nehru Technological University, Hyderabad) Telangana, India.

⁴Professor, CMR College of Pharmacy, (Affiliated to Jawaharlal Nehru Technological University, Hyderabad) Telangana, India.

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*Address for Correspondence

G.S.Sharma

Associate Professor,

Department of Pharmaceutics,

CMR College of Pharmacy,

(Affiliated to Jawaharlal Nehru Technological University, Hyderabad)

Telangana, India.

Email: sharmampharm@gmail.com



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ABSTRACT

Orodispersible tablets (ODTs) are becoming more and more popular among innovative oral drug delivery systems because they enhance patient compliance and offer a few more benefits over other oral formulations. ODTs are solid unit dose forms that quickly disappear or break down in the tongue without requiring water or chewing. In particular, elderly as well as pediatric patients who faces trouble in consuming traditional medications might benefit from the use of ODTs. This review outlines the different ideal properties, selection of drug candidate, types of super disintegrants used, limitations, the disintegrants used, technology created for ODTs. Scientists have used a variety of techniques to create orodispersible tablets. Nonetheless, the compression approach is the most often used way of preparation. The cotton candy process, melt granulation, phase-transition process, sublimation, freezedrying, and quick film formation are other unique techniques. ODTs are also assessed in the areas of hardness, friability, content uniformity, weight uniformity, disintegration test, and dissolution test of solid dosage forms and various applications of oral dispersible tablets.

Keywords: ODTs, Patient compliance, Super disintegrants, Compression technique, Dissolution test.





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INTRODUCTION

Pharmaceutical tablets are characterized as biconvex, flat, or solid plates that are compressed, either with or without diluents, to make a pharmaceutical or drug combination. This is in accordance with the Indian Pharmacopoeia. The quantity of medication and the preferred route of administration have a significant impact on their size, weight, and form. The most popular and advised method of delivering medication, including both liquid and solid dose forms, is orally. ODTs are taken without water and dissolved with saliva. Unless pre-gastric absorption shields it, the active component dissolves quickly in saliva regardless of membrane contact. This study aims to investigate the state of ODT technology at the moment, as well as the ODT characterization and sustainability of drug candidates. Other names for ODTs include orodispersible tablets, fast-acting tablets, mouth dissolving medications."Orodispersible tablet" is the word used by the European Pharmacopoeia to characterize tablets that, after being ingested, dissolve rapidly in the mouth (three minutes). ODT is described as "a solid dosage form containing a medicinal substance or active ingredient which disintegrates rapidly, usually within a matter of seconds, when placed upon the tongue".

PROPERTIES OF ORODISPERSIBLE TABLETS (6, 7)

- It doesn't need water to swallow.
- Dissolves or disintegrates in the oral cavity within less time.
- Show less susceptibility to external factors like temperature and humidity.
- A significant drug loading.
- ODTs must possess flexibility and compatibility with the existing processing and packaging machinery.
- ODTs must also be strong enough to withstand the rigorous demands of manufacturing and post-production processes.

ADVANTAGES OF ORALDISPERSIBLE TABLETS (8, 9, 10)

- ➤ Improved stability
- ➤ Suitable for patients and prescribers, this medication offers greater convenience and compliance for controlled/long-term usage.
- > It promotes patient compliance.
- > Reduces antimicrobial resistance.
- Makes distribution and procurement logistics easier to manage.
- There are the recommended dose form for faster drug administration and higher bioavailability without going through the stomach.
- They may be used to treat a variety of patients, such as the handicapped, travelers, and those who are always on the go and don't always have access to water. They are quite useful.

DISADVANTAGES OF ORALDISPERSIBLE TABLETS (11)

- > It may be necessary to provide the medication more often
- Possibility of dose dumping
- Reduced ability to make precise dosage adjustments
- > Lack of mechanical strength
- ➤ Intervention with rapid pharmacological treatment is not feasible.
- Diminished possibility of precise dosage modification.

SELECTION OF THE DRUGS FOR ODTS (12)

- Medications can diffuse and divide into the GIT top epithelium (log P>1/2); ODT formulations are ideal for those that are efficient to penetrate oral cavity.
- Patients who are taking ach medications at the same time possibly not a suitable candidate for these medications.





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> ODT formulations may not be suitable for patients with reduced salivary flow leading to dry mouth

TYPES OF SUPERDISINTEGRANTS USED (13)

- Cross povidone
- Micro crystalline cellulose
- ➤ Sodium starch glycolate
- Sodium carboxymethyl cellulose or croscarmellose sodium
- Calcium carboxymethyl cellulose

FACTORS CONSIDERED FOR SELECTION OF SUPER DISINTEGRANTS (14)

- ➤ When the medicine is taken orally and then mixed with saliva, it should cause the tablet to dissolve.
- > It should possess a smooth flow since it enhances the overall blend's capacity for flow.
- ➤ It should be compatible enough to generate fewer brittle tablets.

LIMITATIONS OF ORODISPERSIBLE TABLETS (15, 16)

- > When making ODTs, the soluble diluents often give hydroscopic dosages, which can cause stability issues.
- ➤ The tablets are unpleasant to take in the oral cavity if improperly prepared.
- > ODTs are not appropriate for drugs that react to light.

CHALLENGES IN THE FORMULATION OF ODTS (17, 18, 19)

Mechanical strength and disintegration time

Generally speaking, oral dispersible tablets dissolve in less than one minute. There's a greater likelihood that ODTs may break during packing, transit, or patient handling because many of them are brittle. Naturally, the disintegration process will be slowed down by increasing the mechanical strength.

Taste masking

Many medications have an unpleasant taste. Consequently, to stop bitter medications from leaving the tongue feeling tasted, excellent flavor masking is needed.

Size of tablets

The medication ease of use is determined by its size. Tablets bigger than 8 mm are said to be the simplest to manage, and tablets smaller than 7-8 mm are said to be the most easily ingested. That makes it difficult to design a tablet that is portable and manageable in terms of size.

Amount of drug

Generally speaking, the USP advises that the weight of the ODT pill not go beyond 500 mg. Smaller than 60 mg of soluble medication and smaller than 400 mg of insoluble medication should be administered by lyophilized dosage form. This characteristic is especially difficult to work with when creating oral films or wafers that disintegrate rapidly.

Hygroscopicity

In typical temperature and humidity conditions, a number of dosage forms intended for oral dissolution become hygroscopic and lose their physical integrity.

TECHONOLOGIES FOR PREPARING ORODISPERSIBLE TABLETS (20, 21, 22, 23)

- 1. Freeze-drying or lyophilisation
- 2. Melt granulation
- 3. Cotton candy process





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- 4. Phase transition
- 5. Fast dissolving films
- 6. Direct compression.

Freeze drying or lyophilisation

The process of freeze drying or lyophilisation involves removing the solvent from a frozen pharmaceutical suspension or solution that includes excipients to aid in the formation of structures. These pills disintegrate fast due to their porous and generally thin structure. Excipients with a glassy, amorphous porous structure and the medicinal ingredient created by freeze drying both improve solubility.

Melt granulation

Pharmaceutical powders can be efficiently agglomerated using the melt granulation process by using a melt-able binder. This approach has the benefit of not using organic solvents or water, as compared to traditional granulation. To finish this process, high shear mixers are employed. The heat generated by the impeller blades' friction or a heating jacket in these mixers elevates the product temperature over the melting point of the binder.

Cotton candy process

This process is named for the unique spinning mechanism it uses to create crystalline structures that resemble cotton candy and floss. Another name for it is the candy floss approach. In order to create a matrix of polysaccharides, or saccharides, cotton candy is made by spinning and flash melting at the same time. Partially recrystallized matrix is used to improve flow characteristics and compressibility. Subsequently, After the candy floss matrix is pulverized, it is blended with the components for the active as well excipient ingredients, and compressed into ODT. The process results in enhanced mechanical strength and the capacity to manage large dosages of medication.

Phase transition

An innovative method for producing ODTs with the right amount of hardness by utilizing the sugar alcohol phase transition. By compressing and heating two sweeteners and alcohol tablets—one having a significant melting point and the other having a low point of melting—ODTs are produced using this process. Because of the stronger linkages created by heating the particles, the tablets are more durable than they otherwise would have been because of their incompatibility.

Fast dissolving films

It's a novel field of oral dispersible tablets facilitates supplement and drug intake. Medication as well as flavor-masking substances that are permitted to solidify as a film as soon as the solvent disappears. For bitter medications, the film can contain coated drug micro-particles or resin adsorbate. This film quickly melts or dissolves when ingested, releasing the medication in suspension or solution.

Direct compression

ODTs, it can be produced with standard tablet production and packaging equipment and because tableting excipients with better flow, compressibility, and disintegration qualities are readily available. These include sugarbased excipients, effervescent agents, and tablet disintegrants.

Superdisintegrants

Superdisintegrant additions primarily influence the pace of disintegration and, thus, the dissolution in many direct compression-based ODT methods. Additional formulation elements like effervescent agents and water-soluble excipients speed up the disintegration process even further.





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Effervescent agents

The patented Orasolv technology (OT) is based on the evolution of CO2 as a disintegrating mechanism and is widely applied in the development of over-the-counter medications. The product has a mild effervescent quality and contains micro-particles. The effervescent ingredient is activated by saliva, which leads to the tablet disintegrating.

ADVANCEMENTS IN ODT TECHNOLOGIES (24, 25, 26, 27, 28)

- 1. **Zydis technology:** This process encircles the tablet with a rapidly dissolving carrier substance. so that when it gets to the mouth, it will swiftly decompose. To guarantee product stability, these pills include a range of adjuvants. Since the pill was frozen and dried, it is immune to germs. Usually, these pills are kept safe from ambient moisture via blister packs.
- 2. Orasolv technology: CIMA laboratories created this technique. Emulsifying disintegrating agent and flavor masking agent are used to disguise the disagreeable taste of API. To prepare tablets, additional standard equipment is used. To ensure rapid breakdown and softness upon completion, the tablets must to be crushed using a minimal compression force. Furthermore, to maintain the stability of these unpleasant dosage forms over the course of their shelf life, much caution must be used when storing them.
- 3. **Durasolv technology:** The approach was also created by CIMA Labs. The ingredients in composition are the medication, lubricant, and fillers. The same conventional tools used to produce tablets may also be used to make tablets with Durasolv technology. Additionally, these goods don't require specific packing for storage. This method is usually applicable to medications that have a modest quantity of active component.
- 4. Flashtab technology: This technique is yet another cutting-edge way to make oro-dispersible pills. Prographarm labs is the company that created the flashtab technology. Micro crystals are the form of active component utilized in the formulation. The production process can employ a variety of conventional tablet-making methods.
- 5. **Pharma burst technology:** As the name implies, the goal of this technique is to release the drug into the mouth right away. SPI Pharma, New Palace, provided protection for this tactic. This process produces a product that instantly releases the medication from its dose form by combining several specific excipients. One of the unique components utilized to create a powerful, quickly-melting tablet is mold able saccharine.

EVALUATION OF ORODISPERSIBLE TABLETS (29, 30, 31, 32, 33, 34, 35)

Content uniformity

Tablets with a content uniformity test score of fewer than 25 milligrams, or 25% of a tablet, must pass. All 10 random dose units have the same quantity of active component, as determined by the test procedure. A preparation is considered successful if each individual piece of information accounts for 85–15% of the average.

Weight uniformity

Twenty tablets are weighed individually, their average weights are determined, and the weights of each tablet are compared to the average weight in order to perform the weight variation test.

Hardness

Measuring the force required to break a tablet across its diameter yields the tablet's hardness. The hardness of the tablet controls its resilience to abrasion, cracking, and chipping during handling, storage, and preparation for use. ODTs often keep their hardness lower than that of normal tablets since a higher hardness causes the tablet to dissolve more slowly.

Friability test

The weight loss of a tablet within its container due to fine particle removal from the surface is known as friability. Tablets can be tested for friability using the Roche Friabilator.





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Water absorption ratio

Three Folds of tissue paper were put into a small Petri dish that held six milliliters of water. The predicted duration for full soaking was calculated by placing a tablet on the paper. Next, a weight was taken off the moist pill.

Disintegration time

The experiment was conducted on six tablets using the apparatus described in I.P.-1996. Distilled water at $37^{\circ}\text{C} \pm 2^{\circ}\text{C}$ served as the disintegration medium, and the time it took for the tablet to completely disintegrate with no palatable mass remaining in the apparatus was measured in seconds.

Dissolution test

This test is significant because it provides the drug-release profile. They are both USP dissolution test equipment that you may use. Rapid dissolution characterizes orodispersible tablets. Orodispersible tablet fragments or fractured tablet masses can become trapped at the inside top of the spindle of the USP Type I basket device, not withstanding its effectiveness. A poor and ineffective agitating technique might result in an inaccurate dissolving profile. Type II is recommended because of its repeatable-dissolution profile.

Moisture-uptake studies

This study on orodispersible pills is noteworthy. This investigation aims to assess the stability of the pills. The necessary humidity was attained by storing a saturated sodium chloride solution in the bottom of the desiccators for three days. Weighing the pills, we record the weight increase as a percentage.

Packaging

During manufacture and storage, packaging must be handled carefully to preserve the dose of other fast-dissolving dosage forms. Depending on the application and marketing goals, the oral delivery system can be packaged in a single pouch, blister card with multiple units, multiple unit dispenser, or continuous roll dispenser.

APPLICATIONS OF ORALDISPERSIBLE TABLETS (36)

- 1. The goal is to create dose forms that dissolve orally while utilizing already available disintegrants
- 2. To significantly enhance the ODTs' current technology.
- 3. To get ODTs by optimizing the mix of excipients or disintegrates.
- 4. To design and choose appropriate packaging materials and reasonably priced products.

FUTURE PROSPECTIVE FOR ORODISPERSIBLE TABLETS (37)

Many ODT producers face hurdles in the future related to cost reduction through the use of standard equipment, flexible packaging, capacity to hide taste. Orodispersible tablets can be a good option for medications like protein and treatments based on peptides, which are not very bio-available when taken as traditional medicines since they often break down quickly in GIT. Furthermore, the possibility exists to produce controlled release oral drug delivery systems (ODTs) utilizing diverse drug carriers.

CONCLUSION

The most exciting new approaches to medicine delivery is the use of orodispersible tablets. Traditional solid dosage forms may not be as effective as ODTs. ODTs functions like solution inside the body and upon administration, as a solid dose form external to the body. Time to action, bioavailability, convenience, and patient compliance have all increased as a consequence. This dose form should be used carefully because of its low mechanical strength. Future drug production may employ ODT for a great deal more kinds of medications. This medication delivery technique may result in improved patient compliance and ultimately higher clinical output because of the broad relevance of ODT. The development of several new medication classes in the form of ODT is possible.





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REVIEW ARTICLE

Update Review on Iridoid Glycosides: Major Focus on Phytochemical and Pharmacological Versatility

Monika¹ and Neeraj Kumar^{2*}

¹Assistant Professor (Research Scholar), School of Pharmaceutical Sciences, Shri Guru Ram Rai University, Uttarakhand, India.

²Professor, School of Pharmaceutical Sciences, Shri Guru Ram Rai University, Uttarakhand, India.

Received: 29 Dec 2023 Revised: 19 Jan 2024 Accepted: 14 Mar 2024

*Address for Correspondence Neeraj Kumar

Professor,

School of Pharmaceutical Sciences,

Shri Guru Ram Rai University, Uttarakhand, India.

E mail: neeraj_sidana2007@rediffmail.com



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ABSTRACT

The review covers about active substances that found the plant world are called iridoids. Various plants, especially those in the Gentianaceae, Loganiaceae, and Rubiaceous families, contain a group of naturally occurring substances known as iridoids. Numerous biological actions and potential health advantages of these substances are well-known. In traditional medicine and pharmaceutical research, iridoids are of interest because they frequently demonstrate antioxidant, anti-inflammatory, and antibacterial effects.Iridoids have been investigated for their possible function in boosting the immune system, promoting digestive health, and assisting in the management of many chronic conditions. These are the varieties of monoterpenoid, also present in some mammals, in the general form of cyclo-pentanepyran. They come from the biosynthesis of 8oxogeranial and are normally present in plants as glycosides, frequently linked to glucose. Iridoids are bicyclic cis-fused cyclo-pentanepyrans from a structural perspective. Oleuropein and amaro-gnetin are examples of secoiridoids, which are created when a bond in the cyclopentane ring is broken. Iridoids' metabolites are a topic of interest to researchers due to their versatile spectrum of biological activities like neuro defensive, hepatoprotective, anticancer, anti-inflammatory, Anti-Diabetic and lipid lowering activities along with advantages like offering a fresh approach for the creation of new drugs and research focusing. Iridoids have the potential to become new natural medications and supplements because of their bioactive qualities, but additional study is vital to fully comprehend their modes of action and possible uses. In order for a rational consumer of information to be useful in the creation of therapeutic options, the present review's goal is to give extensive information about iridoids and its biological activity.

Keywords: Iridoids, Oleuropein, hypoglycemic, hepatoprotective, hypolipidemic.





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INTRODUCTION

The acetal derivatives of iridodial, which is essentially extracted from the ant's defense secretions, are what make up the group of naturally occurring monoterpenoids known as iridoids. They are generally dicotyledons plants that are more common in the plantae kingdom, including the Scrophulariaceae, Oleaceae, Labiate, Rubiaceous, Gentianaceae, Alliaceae, Loganiaceae, and Verbenaceae families.[1]. These are the components, usually found in the fresh leaves and juvenile stems of plants, but sometimes rarely in the fruits and sprouts, that belong to the broad and extended class of cyclopentane pyran monoterpenes. There are several traditional medications that include iridoids that are used as antipyretics, sedatives, hypotensive agent, bitter tonics, cough medicines, and treatments of wounds and skin conditions. These underlying facts motivated researchers to examine the bioactivities of these phytochemicals. [2,3]. Iridoids possess an ample range of bioactivity like choleretic, neuroprotective, anti-inflammatory, antitumor, antioxidant, cardiovascular, antihepatotoxic, hypolipidemic, hypoglycemic, antispasmodic, antimicrobial, antiviral, immunomodulator, anti-leishmanial, antiallergic and molluscicide.[4] Based on their biosynthesis, chemical and structural characteristics, iridoids have been divided into many subgroups. Four subgroups are distinguished based on the iridoid's structural characteristics: iridoid-glycosides, secoiridoids glycosides, non-glucose iridoids, and bis iridoids. They are naturally more active and include hemiacetal hydroxyl groups. Moreover, they mostly take the shape like glycosides and are link to glucose at the C-1 (-OH) group. Because glycosides have alike qualities like tough ability to modify and facile absorption, they are chosenfor a lead compound. [5,6]. The topic Iridoids have gained attention as an active component that treats a diversity of diseases and disorders as a result of prior studies demonstrating their remarkable biological activity, including hepato-protective, anticancer, hypoglycemic, hyperlipidemic, anti-inflammatory, and effects on various diseases. [7,8] This review's goal is to provide a stronger basis for the creation of novel medicines and research by focusing on iridoids and their biological activity broadly. This will enable future investigation into the chemicals' therapeutic potential.

Main Body Biological Activity of Iridoids Neuroprotective Effects

The body's coordination center, the CNS, acts as the body's remote control for neuronal activity (sending and receiving messages) and movement. Hence, neurological illness results in functional abnormalities in the brain and spine. [9]The causes of neurodegeneration include several elements. Many studies have sought to identify natural medicines that might enhance neuroplasticity in the hopes of avoiding memory loss due to ageing, postponing ageing, and lowering the prevalence of neurodegenerative disorders.[10]

Alzheimer's Disease: Most prevalent type of disorder which is caused by a degenerative brain illness with unclear etiology that often strikes people in their late middle or advanced years. It causes increasing memory loss, poor thinking, disorientation, and changes in personality and mood.[11]. Its disease is indicated by gradual neuronal death, and a deterioration of memory and cognition as a result. The brain's beta amyloid peptide plaques, which produce gliosis, which results in inflammation activation and ultimately cell death, are the primary cause of the gradual cognitive deficiencies.[12]

Mechanism of actions involved

Cholinesterase inhibitions: System acts by inhibiting the acetylcholinesterase enzyme. However, some drugs demonstrated an upsurge in the action of choline acetyltransferase and reversed its depletion, the cause of reduced interneuron connections and shortening of neurite outgrowth. [13]

Improving mitochondrial function: By assisting in mitochondrial dysfunction and the reduction in activity of monoamine oxidase. It also helps in shielding the midbrain neurons from MPTPinduced neurotoxicity.

Filteringdopamine system: Dopamine is a catecholamine neurotransmitter in brain. Iteffects on an increase in DA and DOPAClevel, which helpto slow down Dopamine conversion in nigrostriatal dopaminergic pathway and avert





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the harm of TH-positive cells. The crucial mechanism involved is possibly connected with up-regulating the creation of a glial cell-derived neurotrophic factor in the striatum.[14].

Preventing inflammation:It act byinhibiting microglial cell initiation, the construction of proinflammatory cytokines. Glial cells in the midbrain shows dose-dependent rescued dopamine neurons from chemical induced neurotoxicity. In addition, it also exhibited neuroprotective effects in the D-gal-induced ageing of mice's brains, and its mode of action may be partially explained by the marked decline in inflammatory cytokines. [15-16]

Anti-apoptotic Pathway: By conquering activation of ERK and by monitoring the generation of NO & NOS in rotenone-activated primary cells, it helps in preventing the death of midbrain neurons[17-20]

Parkinson's Disease: Tremor, bradykinesia, and postural instability are symptoms of this slowly progressing neurodegenerative illness that affects the basal ganglia.[21]

In Parkinson disease there is -

- i.) Degeneration of dopaminergic neurons at the fibers nerve endsthat go from the substantia nigra to the striatum causes dopamine to no longer be produced.
- ii.) Characteristic alterations in depigmentation are caused by the loss of the neurons that carry melanin.
- iii.) Lewy bodies' formation (a sticky protein lump that disrupt the normal functions of the brain).
- iv.) Akinesia, stiffness, and bradykinesia are symptoms of dopamine deficiency.[22]

Mechanism of Action of Iridoids in PD

The improved locomotor function by increase in striatal dopamine, TH-positive neurons anddopamine transporter levels. Notably, it raised GDNF protein levels, this elevates GDNF expression is thought to have restored dopaminergic neuron function. [23-26]

Anti –inflammatory Activity

Iridoids' primary health-improving properties are their anti-oxidative and anti-inflammatory properties. Iridoids' ability to suppress proinflammatory cytokines, COX-2,PGE2, and TBX2 accounts for their anti-inflammatory properties. While certain iridoids, such as the hydrolyzed derivative of harpagiferid, are chemically (and physically) comparable to celecoxib, a selective COX-2 inhibitor, and PGE2.In response to inflammation, neutrophils produce a number of hazardous substances, including ROS, that have the natural ability to engulf and kill a variety of bacteria (Reactive oxygen species). Both acute and chronic illnesses have been linked to excessive levels of ROS generated by activated neutrophils through NADPH oxidase. So, it's crucial to prevent inflammatory regions from producing too much ROS from leukocytes.[27-28]One such iridoid is the root of *Gentiana straminea*, which contains the anti-inflammatory agent Loganic acid, which is extensively found in the *Gentianaceae* family. While LA has a high level of COX inhibition and a better affinity for COX-2 than COX-1, it may not have serious adverse effects such an increased risk of gastrointestinal ulcers. Moreover, because of its strong affinity for COX-2. Moreover, it inhibits the production of superoxide in human neutrophils caused by theformyl-methionyl-leucyl-phenylalanine, "a powerful chemotactic factor for human neutrophils".[29]

Hypoglycemic and Hyperlipidemic effect

A series of conditions known together as diabetes are characterized by elevated blood sugar levels. Disease that affects the majority of the people on a daily basis. It is found that Type-2 diabetes is the one that affects the greatest number of individuals. It is a chronic illness that is linked to lifestyle choices and is accompanied by a number of problems.[30]

Pathogenesis of type -2 DM

- Insulin resistance due to Genetic predisposition and Obesity Lifestyle factors.
- dysfunction of islet β cell
- Increase hepatic glucose output
- Decrease peripheral glucose uptake[31]





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Study reveals that decisive gears of the insulin resistance syndrome and criticalissues for heart related disorders are hyperlipidemia and hyperinsulinemia. In various studies, it was revealing that iridoids efficaciously reduce hyperlipidemia and hyperglycemia. It was also discovered that few iridoids play a significant part in a few metabolic diseases. One such iridoid, oleuropein, had a hypoglycemic impact as well as an increase in glucose tolerance *in vivo*. Its mode of action was recently clarified by revealing its potential and positive effects by lowering oxidative stress and boosting the body's own antioxidant defenses. Thus, study reveals a new mechanism of treatment of hypoglycemic effect through antioxidant action [32]. However, Iridoidsseparated from the leaves of *G. Jasminoides* (Rubiaceous) shows that deacetylasperulosidic acid methyl ester had hypoglycemic effect and decreased blood glucose levels in normal mice.[33-34]

Mechanism of Action of hypoglycemia[35]

- Insulin stimulus
- Increase and Improve insulin sensitivity andresistance
- Restoration of pancreas
- Action on metabolism of glucose
- Regulating flora of intestine

Hepatoprotective Activity

The liver, the body's biggest and most important organ, is responsible for a number of vital biological processes, including metabolism, detoxification, the synthesis of proteins, and the production of biochemicals required for development and digesting. The organ's function exposes it to harm from many types of metabolic products, toxins, endotoxins, viruses, or medications. [36]Oxidative stress is the main factor to take into for liver disease. Damage to the liver is mostly caused by increase in the reactive oxygen or nitrogen and also in decrease in antioxidant molecules. It is depleted by highly reactive metabolites. Glutathione is required for the cleansing and to remove toxins of glutathione-S-transferase and also leads to causes cells to die. [37] Antioxidant is the function to perform in this situation. Iridoids take the lead thanks to their antioxidant capacity. The activity is both direct and indirect, by removing reactive oxygen species and by activating the anti-oxidant defenses within the body. [38-40] Iridoid, guards the liver against steatosis, fibrosis, and necrosis. According to various studies, they lower liver/body weight ratio, bilirubin, ALT, AST, and ALP. Iridoids also limits the fibrosis process by decreasing the accumulation of collagen in the liver and the inflammation by loweringIL-1, TNF, IL-18, IL-6 and COX-2 in a dosedependent way. On the basis of above result a conclusion is drawn that iridoids' hepatoprotective effect are because of their facilitation of drug metabolism, enhancement of mitochondrial dysfunction, decrease in oxidative stress and by suppression mitogen activated protein kinase signaling pathway. [41]

Antioxidant Activity

Each and every cell in the body depends on oxygen. Cells cannot produce energy without oxygen; therefore, metabolism is less efficient. Cellular functions such controlling gene expression, signal transmission, cell growth and development and cell death depend on it for survival. The body needs oxygen for oxidative phosphorylation because it facilitates the aerobic metabolism of glucose, which produces the energy needed by the body. Then, the formation of reactive oxygen species occurs. Now, in order to adapt to the environment, the body develops a self-antioxidant defense system, which includes vitamins, coenzyme Q, lipoic acid, glutathione (GSH), and other substances, all of which work together to prevent ROS from being produced. ROS can be eliminated even by antioxidant enzymes. In a typical physiological circumstance, there is a continual fluctuation in the creation and removal of ROS. Although it triggers an event that results in an excess of ROS and a lack of antioxidants, which leads to oxidative stress, when subjected to detrimental external stimuli.[42]This oxidative stress causes cell necrosis and apoptosis by destroying the structure and functionality of several macromolecules. The research study shows thatan iridoid(Aucubin) is a potent antioxidant. It either directly lowers levels of ROS, Malondialdehyde, and 4-hydroxynonenal, as well as efficiently remove oxygen, free hydroxy and 1,1-diphenyl-2-picrylhydrazyl radicals, or it does the opposite by raising the antioxidative activities of Superoxide dismutase, catalase, andGlutathione peroxidase. By balancing





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oxidation and the antioxidant defense system better, therefore iridoids shows anincreased antioxidant ability to prevent damage to the gastric mucosa, endothelium, cardiac remodeling, and diabetic encephalopathy.[43-45]

Anti-tumor Activity

A tumor is an atypical mass of tissue that progresses as result of excessive cell growth, division, and survival rather than demise. Both benign and malignant tumors exist. It is among the world's major causes of death. Studies reveal that Gentipicroside, Valjatrate E, and Sertiamarin B are a few iridoids that do have anti-tumor effect. [46]. Gentiopicroside has been reported to show antitumor activity against ovarian cancer cell line (SKOV3). Its anticancer effects were determined using Methyl Thiazolyl Tetrazolium, Apoptosis by 4', 6-diamidino-2-phenylindole annexin V or propidium iodide (PI) double staining, Mitochondrial membrane potential (MMP) and protein expression by western blotting. [47] All resultsuggests that gentiopicroside may prove to be an effective lead chemical in the treatment and management of ovarian cancer. [48]

The Sertiamarin B, a new seco-iridoid, a chemical component which is identified from *Swertia mussotii*'s aerial part. Four human tumor cell lines (HCT-116, HepG2, MGC-803 and A549)exposed to it showed anti-tumor action. It demonstrated substantial cytotoxic activity against the MGC-803 cell line (IC50 = 3.61), and its ability to efficiently suppress cell growth raises the possibility that it might serve as a novel gastric cancer treatment agent. [49]

Anti-bacterial Activity

According to studies on iridoids various researchers demonstrated that they have antibacterial properties. Isoplumericin, plumericin, galioside, and gardenoside are a few of them; they have all been linked to anti-bacterial action. An intriguing action on *Staphylococcus aureus* was discovered when an iridoid (seco-iridoid glycoside)gentipicroside showed significant efficacy (MIC 6.3103 mg/ml). Likewise, Sweroside (an iridoid) isolated from the rootstock of *Scabiosa columbria* (Dipsacaceae) was used against *B. cereus, B. pumulus, B. subtilis, M. kristinae, S. aureus, E. coli, K. pneumoniae, P. aerugenosa, E. cloacae* and it showed good antibacterial activity. A group of various non-glycosidic iridoids likemussaenin A, gardendiol, isoboonein, and rehmaglutin D showed a potent anti-bacterial action. It shows that all glycosidic and non-glycosidic iridoids have antibacterial action similar to chloramphenicol.[50]

CONCLUSION

Iridoids are a substance that is extensively dispersed across the plant world; as previously indicated, it is particularly common among dicotyledons plants. Research in contemporary pharmacology have demonstrated their biological and pharmacological activity, high therapeutic value, and usefulness in the treatment of tumors, hepatoprotection, antimicrobial, neuroprotection, antioxidant, hyperlipidemia, and inflammation. Many biological functions of iridoids were investigated in this review paper. Iridoids' kinds, structure, and how the placement of different functional groups alters how active they are all discussed. Oleuropein, *C. officinalis*, and *G. jasminoides* are used as hypoglycemic agents. Catalpol and picroliv are used for hepatoprotective activity. Aucubin is used as an antioxidant. Gentiopicroside and Sertiamarin B are used in anti-tumor activity. Aucubin and Sweroside are used as antioxidants. Iridoids like catalpol are used for neurodegenerative diseases like Parkinson. This shows that the iridoids' versatility and current study on their activity have been important in luring scientists to learn more about iridoids in order to unlock these chemicals' untapped potential. This leads us to the conclusion that the iridoids group may prove to be a medicine with significant therapeutic potential and that more research into this area may lead to the creation of treatments with greater efficacy.

Abbreviations

CNS Central Nervous System

MPTP 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine

DA Dopamine





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DOPAC 3,4-Dihydroxyphenylacetic acid

ERK Extracellular signal-regulated kinase

PD Parkinson's Disease

GDNF Glial cell-derived neurotrophic factor

COX Cyclooxygenase
PGE2 Prostaglandin E2
TBX2 Thromboxane-B2
ROS Reactive oxygen species

DM Diabetes Mellitus

ALT Alanine aminotransferase AST Aspartate aminotransferase ALP Alkaline phosphatase

DPPH 1,1-diphenyl-2-picrylhydrazyl

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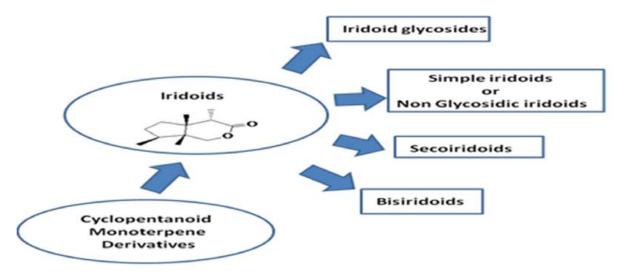


Fig1.: Chemistry and classification of iridoids.





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REVIEW ARTICLE

Application of Nanotechnology in Healthcare Sector: A Perspective Review

V.Sireesha^{1*}, N. Chaitanya², K. Megana² and Meghana.Ch²

¹Assistant Professor, CMR College of Pharmacy, Medchal, (Affiliated to Jawaharlal Nehru Technological University), Hyderabad, Telangana, India.

²Pharm D Student, CMR College of Pharmacy, Medchal, (Affiliated to Jawaharlal Nehru Technological University), Hyderabad, Telangana, India.

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*Address for Correspondence

V. Sireesha,

Assistant Professor,

CMR College of Pharmacy,

Medchal, (Affiliated to Jawaharlal Nehru Technological University),

Hyderabad, Telangana, India.

Email: sireeshaganesh59@gmail.com,



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ABSTRACT

Nanotechnology is a fast-developing area as nanomedicine in engineering and medicine that holds the key for improving medical care in many areas. Nanoparticles have possible applications in diagnosis, targeted medicine products, pharmaceutical products, and biomedical implants. A detailed discussion of the use of nanotechnology in different sectors, including medicine and health, Drug delivery, for treating cancer and cardiovascular disease and diagnosis. Our article discusses about the use of different nanotubes in the delivery of drugs for the treatment of cancer. This article explains how nanotechnology is used in dentistry, nutrition, Covid vaccinations, and HIV/AIDS. Thus our Perspective review gives brief and well organized review on Nanotechnology that should be valuable to researchers, engineers and scientists for future research project.

Keywords: Nanotechnology, nanodevices, nanometer scale, Nanomedicine, Nanomatrials, Polyethylene glycol

INTRODUCTION

The idea and concept of Nanotechnology was introduced by the American physicist and Nobel Prize laureate Richard Feynman in 1959. Nanotechnology is the branch of science that manipulates matter on a scale close to the atomic level to create new materials, structures that function well at the Nanoscale. The development and advancement of numerous scientific fields, including medical, industry, energy, materials science and engineering, depend on this technology [1]. A typical dimensional range for nanoscale constructions is between 1 and 100 nano





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meters. Nanomaterials can be produced as atomic-scale structural Manipulation is possible [2]. So these Nanomaterials are unique as they provide large surface area to volume ratio. Utilizing materials with distinctive features at the nanoscale is known as Nanotechnology. The use of Nanomaterials can be found in sunscreen, cosmetics, sporting goods, tyres, electronics and several other items. The use of Nanomaterials in every day products can be divided into 2 types:

- a. Nanomaterials can be merged or added to pre-existing product to improve the performance.
- b. Nanomaterials such as Nanocrystals and Nanoparticles can be used directly to create advanced and powerful devices[3].

Types of Nanoparticles

Several Nanoparticles and Nanomaterials have been investigated and approved for clinical use. These are tiny macromolecules used in medicinal Nanotechnology. Nanomaterials and Nanodevices are the two main categories of nanotools used in pharmaceutical research. Nanoparticles, dendrimers, micelles, drug conjugates, metallic Nanoparticles, etc. make up Nanostructures[4]. Some common types of Nanoparticles are discussed below

Micelles

These are amphiphilic surfactant molecules formed by combination of Lipids and other amphiphilic molecules. Hydrophobic therapeutic drugs can be included into micellar spherical vesicles, which spontaneously aggregate and self-assemble into a hydrophilic outer monolayer and a hydrophobic core in water conditions. As the special qualities of micelles, hydrophobic medications are made more soluble, increasing their bioavailability. Micelles have diameters ranging from 10 to 100 nm. Micelles can be used as therapeutic, imaging, contrast, and drug delivery agents, among other things[5]

Liposomes

Liposomes are spherical vesicles made of lipid bilayers that range in size from 30 nm to several microns. Hydrophilic therapeutic compounds can be incorporated into the liposomal membrane layer of liposomes, while hydrophobic agents can be incorporated into the aqueous phase [6]. One of the top nanosystems being developed for targeted medication delivery to treat different malignancies and cardiovascular illnesses is the nano-liposome. Doxil is the first FDA approved nano-liposome for treating breast cancer, it enhances the effective drug concentration in malignant effusion without the need to increase overall dose [7].

Dendrimers

Dendrimers are macromolecules that have an external functional group and branched repeating units that extend from a central core [8]. Because therapeutic substances can be bonded to the surface groups or enclosed within the internal space of dendrimers, dendrimers are extremely biodegradable and bioavailable [9]. Dendriplexes, which are polyamidoaminedendrimer-DNA complexes, have been studied as potential vectors for delivering drugs, and they show promise in promoting targeted drug delivery, successive gene expression, and enhanced medication efficacy [10]. Because of their transformable characteristics, dendrimers are a potential class of particulate systems for biological applications, including drug administration and imaging [11].

Carbon nanotubes

Carbon nanotubes are spherical molecules made of graphene, a single layer of carbon atoms wrapped up into sheets [12]. Carbon nanotubes can reach remarkably high loading capacities as drug carriers because of their large exterior surface area. Additionally, carbon tubes are attractive as biological sensors and imaging contrast agents due to their distinct optical, mechanical, and electrical properties [13].

Metallic nanoparticles

Gold and iron oxide are examples of metallic nanoparticles. Hydrophilic polymers like PEG with a magnetic core (4-5 nm) make up iron oxide nanoparticles[14]. Metallic nanoparticles have been applied as drug delivery vehicles, optical biosensors, laser-based therapy, and imaging contrast agents[15].





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Quantum dots

Quantum dots (QDs) are fluorescent semiconductor nanocrystals that range in size from 1 to 100 nm. They have demonstrated promise for usage in a number of biomedical applications, including cellular imaging and drug administration. Quantum dots are used in medical imaging because of their unique optical characteristics, small size, high brightness, and stability [16].

Nano-Tube

The identification of different volatile organic chemicals in breath samples from patients with stomach and lung cancer, respectively, has been done using silicon nanowires and carbon nanotubes[17].

Nanobots

Robots of a nanometre (109 m) scale are also known as nano robotics, and they have been used in healthcare, early cancer diagnosis, targeted medication administration for treatment, and pharmacokinetic monitoring of diabetes. Nanobotsdentrifices (dentifrobots), for example, can cover all subgingival surfaces when used as toothpaste or mouthwash, metabolizing any organic material trapped there into safe and odorless fumes. Using correctly configured dentifrobots, pathogenic bacteria that are present in dental plaque are found and eliminated.

Nanowires

Carbon nanotubes, metal oxides, or silicon can be used to create nanowires (NW), which are nanosized channels that permit the passage of electrical current at very low amplitudes. Because of their tiny size and tiny diameter, which is typically 10 nm, they are extremely sensitive to even the smallest change in electrical characteristics. In order to use antibodies as detectors, one option is to attach them to the surface of nanowires. When the antibodies engage with the biomolecules of the target, this causes a conformational change that is detected as an electrical signal on the nanowire. As a result, when multiple nanowires with various antibodies attached are combined into a single device, they can be used as detectors for illnesses like cancer.

Application of Nanotechnology

Nanotechnology have a significant impact in almost all industries. Nanomaterials are directly used to create advanced and powerful devices. Nanomaterials serve as a beneficial use in sunscreen, cosmetic, sporting goods, tyres, electronics^[19]. Additionally Nanotechnologies have revolutionized advances in medicine, specifically in diagnostic methods, imaging and drug delivery. The different fields that find potential applications of nanotechnology are as follows:

Nanotechnology in health and medicine

Nanomedicine is the branch of Nanotechnology and is used for the application of nanotechnology to the fields of medicine and healthcare [20]. It has been utilized to treat some of the most prevalent illnesses, such as cancer, musculoskeletal, psychiatric, neurodegenerative and cardiovascular conditions [21]. Early disease detection and prevention, better disease diagnosis, appropriate therapy, and follow-up are all made feasible by nanomedicine. Nanomaterials and nano electrical biosensors are used in nanomedicine. Nanomaterials and the use of nanotechnology allows for the replication or repair of damaged tissue. Tissue engineering, which could revolutionize organ transplantation or the use of artificial implants, uses so-called artificially activated cells. Carbon nanotubes can be used to create advanced biosensors with unique properties. The development of sensors for cancer diagnosis is also being done using this technology. In order to track and image stem cells, promote their differentiation into particular cell lineages, and eventually comprehend their biology, nanotechnology can be a useful tool[22].

In Drug delivery

Nanoparticles are employed in nanotechnology to deliver drugs to precise sites. As the active substance is exclusively deposited in the morbid zone, this procedure uses the required drug dose and considerably reduces adverse effects. Targeted medication delivery has seen a sharp rise in the use of nanotechnology in recent years, particularly in the medical fields. To transport medications, heat, light, or other things, to particular types of cells





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such as cancer cells, nanotechnology is currently being explored. One of the top nanosystems being developed for targeted medication delivery to treat different malignancies and cardiovascular illnesses is the nano-liposome. Drug delivery systems should be carefully designed, using lipid- and polymer-based nano-particles to enhance the pharmacological and therapeutic effects of medications[23]. Thus Nanoparticles find their use in medication encapsulation, block co-polymer-derived micelles, they deliver tiny medication molecules where they are needed. For the active release of medications, nano electromechanical systems are also used. A tailored medication lowers drug consumption and medical costs, making patient care more affordable for medications made with nanotechnology[24].

Need for Nanotechnology in drug delivery

- 1. Side effects can be reduced significantly.
- 2. Drug efficiency is ensured by specifically targeting and killing harmful and cancerous cells.
- 3. Minimised irritant reactions and improved penetration within the body due to their small size.
- 4. Improved drug bioavailability.
- 5. Provide sustained release/controlled release/prolonged release of the drug.
- 6. Tissue damage can be prevented by administering under regulated conditions.
- 7. Nanoparticles are used to administer the medication mare effectively for treatment of head and neck cancer.

For example: Doxorubicin which is highly toxic can be delivered directly to tumour cells using liposomes without effecting heart or kidney[25].

Diagnosis and Imaging

The greatest contribution of nanotechnologies to medicine has been the development of more potent contrast agents for nearly all imaging models. This is because nanomaterials have improved permeability and retention effects in tissues, as well as lower toxicity. The development of nano-enabled imaging diagnostic tools and nano-sensors is a focus of nanotechnology in biomedical engineering. Iron oxide (Fe2O3) nanoparticles have been used as contrast agents to improve magnetic resonance imaging for many years, Approved by US FDA The management of disease prognosis, which is important in determining the survival rate of patients, is enhanced by this. Nanoparticles are being used for molecular and biochemical sensing in cancer research, To find infections on-site, many different nanoparticles have been utilized. One of the most popular nanoparticles for rapid diagnostics is gold nanoparticle (AuNP)[26]. Nanodevices have been investigated for the detection of blood biomarkers and toxicity to adjacent healthy tissues. These biomarkers include tumour-shedding exosomes, circulating tumour nucleic acids, related proteins or cell surface proteins, circulating tumour nucleic acids, and cancer-associated circulating tumour cells. These biomarkers not only aid in the early detection of cancer, which is well recognized, but also in the monitoring of treatment and recurrence. Nano-enabled sensors allow high sensitivity, specificity, and multiplexed readings. Next-generation technology combines capture with genetic analysis to shed more light on a problem[27,28].

In cancer diagnosis and tumour hunting

Nanotechnology provides a speedier and more accurate initial diagnosis in addition to a continuous evaluation of cancer patient care through the use of novel molecular contrast fluids and materials[29]. Chemical modifications of surface of Nanoparticles improve targeted delivery. One of the best modification is incorporating PEG, which avoids detection of Nanoparticle as foreign objects by the body's immune system, thus allowing them to circulate in blood stream until they reach tumour.

For **example**: application of hydrogel in breast cancer is the innovative technology and the Herceptin is a type of monoclonal antibody. To meet the requirement for nutrients and oxygen for cancer cell survival and growth, solid tumours frequently develop an aberrant, excessive vasculature. Therefore, blocking tumour vasculature is a promising therapeutic approach to starve tumour cells. It is essential, nevertheless, that such a therapy be able to recognize and block tumour blood arteries while ignoring those in healthy tissues. Design for a tumour-hunting





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nanorobot that incorporates DNA origami and thrombin in a capsule. In animal models, the DNA robots specifically cause intravascular thrombosis (blood clotting) in tumours, causing necrosis and inhibiting tumour growth[30].

In treating Cardiovascular disease

Another area where the characteristics of nanoparticles may be used is in the treatment of cardiovascular disorders. The main cause of mortality worldwide is cardiovascular disease, and sedentary lifestyles are to blame for the worrisome rise in death rates^[31]. permanent disabilities are caused by these illnesses. Novel therapeutic and diagnostic approaches for the treatment of cardiovascular disorders are made possible by nanotechnologies.Novel formulation of block copolymer micelles are made using PEG and poly(propylene sulphide) this supresses the level of cytokines which helps in managing atherosclerosis [32].

Nutrition

Many antioxidant nutrients and food ingredients lose some of their potency while traveling to the target tissues. Nano-concepts can be used in nutrient delivery to improve everyone's health and fitness, not only athletes, especially when it comes to weight control. Nanotechnology can be used to study microorganisms and change the flavour, colour, and quality of food. These ideas can help advance the research of physiology, nutrition metabolism, and dietary problems[33,34]

Dentistry

Nano-dentistry is a contemporary field in development that has the potential to use mechanical nana-robotic dentifrices for oral health care. This can be used in conjunction with local anaesthetic to treat hypersensitive tooth conditions and perform orthodontic corrections. The use of Silver Nanofluoride (NSF) in newborn dental care has also been demonstrated to be effective by recent research [35].

Energy and Environment

Fuel cells made of carbon nanotubes are used to power cars because they can store hydrogen. Photovoltaic systems use nanotechnology to become more affordable, lightweight, and effective. These systems can also reduce engine pollution through the use of nanoporous filters and mechanically clean exhaust with the aid of catalytic converters made of nanoscale noble metal particles, catalytic coatings for cylinder walls, and catalytic nanoparticles added to fuel[36,37].

Risks Associated with nanotechnology

While nanotechnology has generated a lot of interest in the general public, there have also been a lot of discussions about the safety of nanotechnologies and potential health dangers linked with their use. The use of nanomaterials presents new difficulties, particularly in terms of foreseeing, comprehending, and controlling any possible health hazards. Low-solubility nanoparticles are more hazardous and toxic on a mass by mass basis than larger particles[38]. The body can absorb nanoparticles through the skin, by food, inhalation, or injection during medical operations. Due to their great mobility, nanoparticles may be able to pass through the blood-brain barrier after they have entered the body. Through phagocyte overpopulation and nanoparticles may impact the immune system of the organism. Stress and inflammation responses could be set off, weakening defenses against more dangerous obstacles[39,40]. Significant energy requirements for nanoparticle production and may damage the environment by releasing hazardous and enduring nanoparticles into it. Low rates of recovery and limited possibility for recycling[41].

CONCLUSION

Nanotechnology emerges as a groundbreaking force with transformative potential across multiple sectors, especially in healthcare. Nanoparticles' versatility in drug delivery, imaging diagnostics, and therapeutic applications promises revolutionary advancements. However, challenges regarding safety and environmental impact necessitate ongoing





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exploration. Overall, this comprehensive review underscores nanotechnology's immense promise while highlighting the need for responsible integration and further research for its sustainable use and maximal benefit.

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RESEARCH ARTICLE

Optimization and Partial Purification of Protease from Bacillus Subtilis **AD20 Isolated from Dairy Effluent**

Deepika Jothinathan^{1*}, Anjali Kumari¹, Lavanyasri Rathinavel²

¹Department of Biotechnology, SRM Arts and Science College, Kattankulathur, (Affiliated to University of Madras) India.

²Department of Biotechnology, Mercy College, (Affiliated to Calicut University) India.

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*Address for Correspondence Deepika Jothinathan

Department of Biotechnology, SRM Arts and Science College, Kattankulathur, (Affiliated to University of Madras) India.

Email: deepikabt@srmasc.ac.in



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ABSTRACT

Among the total worldwide enzyme market, 40% of the total protease is produced by microbial protease. The study aims to produce, purify and determine the properties of the protease from a bacterium isolated from diary effluent. The dairy effluent sample was collected from the dairy industry and bacteria were isolated by serial dilution on skim milk agar. The isolated bacteria were screened on skim milk agar for proteolytic activity. Based on the results of primary screening, different bacterial species were separated from the dairy effluent sample. The bacterial strain with the greater zone of clearance was selected for protease production in complex media. The protease positive bacterium was determined by 16S rRNA sequence analysis as Bacillus subtilis AD20. The optimisation studies with different parameters were investigated to produce the maximum extracellular protease. This enzyme was then partially purified with the help of 80% ammonium sulfate precipitation tailed by dialysis. The partially purified protease was studied by SDS- PAGE and Native-PAGE with Gelatin for their molecular weight determination and protease activity respectively. The findings of the characterization investigations indicated that the protease was most active at 55°C and an acidic pH of 4. Among the tested substrates higher activity was evident with casein. In the presence of benzene, protease showed higher activity. According to results obtained it was understood that Bacillus subtilis AD20 produces protease that is temperature resistant i.e., thermostable and acidic. The comparison of enzyme activity with commercial detergent activity showed positive results. The washing performance studies were done which cleared the blood and dal stains.

Keywords: 16S rRNA, Bacillus subtilis AD20, Ammonium sulfate precipitation, Dialysis, SDS- PAGE and Native-PAGE





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INTRODUCTION

Proteolysis, the process by which proteins are broken down into smaller polypeptides or individual amino acids, is catalyzed by the enzyme protease. These enzymes use a process called hydrolysis, in which water breaks peptide bonds, to cleave the bonds within proteins. Proteases are engaged in a variety of biological processes, such as cell signaling, protein catabolism, and the digestion of consumed or ingested proteins [1]. There are proteases in every kind of life. Proteases have undergone numerous independent evolutionary cycles and can catalyze the same reaction using entirely different catalytic pathways. Therefore, despite the fact that proteases are widely distributed, rapid growth, low cultivation area requirements and simplicity of genetic manipulation to produce new enzymes with altered properties that are desirable for their many uses, microorganisms are the preferred source of these enzymes [2]. Due to its high water use, the dairy industry is one of the most polluting in the food business. India's dairy business is predicted to expand quickly in light of the country's rising milk consumption. The waste production and associated environmental problems are also assumed to be amplified [3]. Dairy is one of the major industries causing water pollution. Many processes in the dairy manufacturing might contain pasteurization, cheese, cream, milk powder, etc. Large volumes of milk are handled by the dairy sector, and water is the main processing byproduct. Significant levels of organic milk products and minerals can be found in the milk's extracted water. Additionally, the plant's cleaning process produces caustic wastewater [4].

MATERIALS AND METHODS

Effluent collection

The positive strains for protease were isolated from dairy effluent which was collected from dairy industry, Chennai, Tamil Nadu.

Screening of Bacterial Strains for Protease Production

The protease producing bacterial strains were isolated from dairy effluent. Serial dilution was performed for the dairy effluent sample. The 10⁻⁷, 10⁻⁸, and 10⁻⁹ serially diluted samples were used for the spread plate technique in skim milk agar plates. The plates after overnight incubation, was observed for clear zone formation. A greater zone of clearance was used as a positive strain for further studies by separately inoculating the different stains in skim milk agar. The positive culture was preserved on sterile nutrient medium by quadrant streak plates and slants at 4°C.

Identification of protease-producing by 16S rRNA sequencing

The microbial positive strains were characterized by 16S rRNA using sanger sequencing was performed in Elango Genetics Pvt Ltd.

Optimization of Culture Conditions for Protease Production

The optimal conditions for the growth of the positive cultures and growth curves were determined with different parameters (time durations, pH, carbon source, nitrogen source, and specific substrates).

Effect of Time Duration

The production medium was prepared in a flask and sterilized. The positive culture was inoculated and retained in a shaker incubator at 100 rpm at 37° C. Flasks were kept for incubation in a shaker incubator for 100 rpm at room temperature. The protein content and protease activity were recorded for every 24 hours for 92 hours (i.e., 4 days).

Effect of Different pH

The production medium was arranged with various pH i.e., 5.0, 6.0, 7.0, 8.0 and 9.0. After sterilization, the positive strain was inoculated and retained in a shaker incubator at 100 rpm at room temperature. The protein content and protease activity were recorded every 24 hours for 92 hours (i.e., 4 days).





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Effect of Various Carbon Sources

The production medium was prepared with different 1% carbon sources i.e., starch, dextrose, sucrose, maltose, and fructose and inoculated a loop of positive strain in the media after sterilization. Then the flasks were incubated in a shaker incubator at 100 rpm at room temperature. The protein content and protease activity were recorded every 24 hours for 92 hours (i.e., 4 days).

Effect of Different Nitrogen Sources

The production media was prepared with different 1% different N₂ sources i.e., ammonium acetate, urea, peptone, casein, and yeast extract and inoculated a loop of positive strain in the media after sterilization. Then flasks were maintained in the same rpm and temperature used in the previous experiments. The protein content and protease activity were recorded every 24 hours for 92 hours (i.e., 4 days).

Effect of Various Substrates

The production medium was prepared with different 1 % substrates i.e., green gram husk, rice bran, sugarcane bagasse, grape waste, and kitchen waste and inoculated a loop of positive strain in the media after sterilization. The same conditions were maintained throughout the experiment. The protein content and enzyme activity were recorded every 24 hours for 92 hrs. (i.e., 4 days).

Estimation of Enzymatic Reaction

Using tyrosine (0.2–1.0 mg/L) as the standard, the protease activity method was used to determine the protease activity of the samples. A standard calibration curve was created using the response to the absorbance values of tyrosine standards that were developed in a range of concentrations. A UV Visible spectrophotometer was used to analyze the samples. The assay mixture was incubated for 10 minutes at 55° C in the water bath to measure the protease activity. 100 μ l of 0.5% (w/v) casein in 50 mM Tris-HCl pH 7.2 (substrate) was added to 100 μ l of enzyme solution (supernatant). To stop the process, 100 μ l of 15% TCA was added in autoclaved distilled water to the enzyme-substrate solution. After 10 minutes in an ice bath, the mixture was centrifuged for 10 minutes at ambient temperature at 10,000 rpm. The pellet was then removed. 200 μ l supernatant was taken in fresh micro fuge tube, then 1 ml NaOH (1N) and 200 μ l Folin & Ciocalteu's Phenol reagent was added [5]. The control sample had the similar composition excluding the enzyme. The enzyme activity was documented for every parameter by UV Visible spectrophotometer at 660 nm after 1 hour incubation at room temperature.

Quantification of Protein Concentration

Using bovine serum albumin (BSA, 0.1 mg/ml) as the standard, the Bradford method was used to calculate the total protein concentration of a sample [6]. Aside from the enzyme, the makeup of the control sample was the same. The standard calibration curve was created in response to the absorbance values of the BSA standards, which were manufactured in various concentrations. The absorbance was measured in a UV visible spectrophotometer at 595 nm.

Mass Production of Crude Enzyme Extract

2 ml overnight culture was inoculated in 1-liter sterilized complex media and kept in a shaker at 100 rpm at 37°C for 3 days. Based on the previous experiments, the parameters that produced the maximum protease activity of the organism were chosen for mass production. To get the culture supernatant, the bacterial culture was centrifuged for 20 minutes at 5000 rpm and 4°C . After transferring the culture supernatant into the sterile flask, the pellet containing the cell debris was taken out.

Partial Purification of Enzyme

Ammonium sulfate was dissolved to 80% saturation (w/v) in the crude extracellular enzyme extract [7]. The ammonium sulfate has been gradually added into the culture filtrate by continuously stirring in a cold room. After dissolving, the flask was kept for overnight incubation for the precipitate formation of salt and protein at 4°C. After





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collecting the precipitate using centrifugation at 10,000 rpm and 4° C for 10 minutes, the pellet was reconstituted in a brown bottle containing 50 ml of 50 mM Tris-HCl buffer at pH 7.2.

Preparation of Dialysis Membrane

The dialysis membrane was washed for 5 minutes in hot water. The enzyme solution was loaded in the dialysis membrane - 70 i.e., DEAE Cellulose column (width: 29.31 mm; diameter: 17.5 mm) which was dissolved with 50 mM Tris HCl buffer of pH 7.2. This membrane was dipped in 20 mM Tris HCl buffer (pH 7.2) for 6 hours and repeated for 2-3 times by changing the 20 mM tris HCl buffer.

Electrophoretic Studies

Sodium Dodecyl Sulphate-Polyacrylamide Gel Electrophoresis (SDS-PAGE)

The SDS-PAGE procedure was performed using Laemml's 1970 method [8]. After five minutes of heating the samples in boiling water at 100 °C, the molecular mass of protease was ascertained using separating gel with 10% acrylamide concentration and stacking gel with 5% acrylamide concentration. The electrophoresis was conducted after loading 25 μ l of the sample-sample buffer mixture and 10 μ l of the molecular weight marker onto the gel. The gel was passed through a resolving gel at 100 V and a stacking gel at 50 V for duration of 20 minutes. Coomassie staining was applied to the SDS-polyacrylamide gel for 30 minutes following the electrophoretic run. Finally, bands were evident after the gel was destained with the destaining solution and incubated for 12 hours.

Native Polyacrylamide Gel Electrophoresis (Native-PAGE)

The protease enzyme's approximate size was discovered using native PAGE. 0.1% gelatin was present in the polyacrylamide gel that was cast using 15% resolving gel. Using a typical gel loading buffer, the sample was loaded onto the gel. Due to the breakdown of gelatin in that area, a clearing was observed at the band responsible for protease activity when the enzyme sample was run on a polyacrylamide gel containing gelatin [9]. Under natural circumstances, the concentrated crude protease was electrophoresed [10]. Following electrophoresis, the gel was ran through three rounds of washing in 0.1% Tween 80 for thirty minutes each, and then it was cleaned with sterile distilled water. During two hours, the gel was immersed in 100 mM Tris HCl, pH 8.0, with 5 mM CaCl₂.

Characterization Studies of Protease

Impact of pH on Protease Activity and Stability

Protease activity was measured using a 50 mM Tris-HCl buffer at various pH levels for ten minutes at 55° C. Using conventional test settings, the optimal pH of the protease was ascertained by measuring its activity in the presence of buffers at various pH values (5.0, 5.0, 6.0, 7.0, 8.0, 9.0, and 10.0). Using spectrophotometry, the change in absorbance at 660 nm was recorded, and the relative activity was calculated.

Impact of Temperature on Protease Activity and Stability

The impact of varying temperatures on protease activity was examined through the use of 50 mM Tris-HCl buffer. Under normal assay environment, the optimal temperature of the protease was ascertained by measuring its activity at various temperatures (4, 25, 37, and 55° C). Prior to the test and for ten minutes throughout the test, the protease and the particular substrate, casein, were incubated at the appropriate temperatures. Using spectrophotometry, the change in absorbance at 660 nm was recorded, and the relative activity was calculated.

Substrate Specificity of Enzyme

By monitoring the enzyme's activity toward various substrates (BSA, casein, peptone, tryptone, and gelatin), the substrate specificity of the enzyme was ascertained. These various substrates were prepared in 50 mM Tris-HCl buffer at concentrations of 0.5% (w/v) according to normal assay conditions and the enzyme's activity was assessed by incubating them for 10 minutes at 55° C. Using spectrophotometry, the change in absorbance at 660 nm was recorded, and the relative activity was calculated.





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Organic Solvents of Enzyme

The organic solvent specificity of the enzyme was estimated by measuring activity towards different solvents (Ethanol, Methanol, Benzene, and Dimethyl Sulfoxide). Using these various organic solvents, which were produced in 50 mM Tris-HCl buffer at solvent concentrations of 10% under standard test conditions, the enzyme activity was determined and the mixture was incubated for 10 minutes at 55° C. Using spectrophotometry, the change in absorbance at 660 nm was recorded, and the relative activity was calculated.

Analytical Methods of Partially Purified Protease by Skim Milk Assay

Protease activity was analyzed using skim milk 0.5% (w/v) with nutrient agar plates. Five wells were created, control (10 μ l), 20 μ l, 30 μ l, 40 μ l, and 50 μ l partially purified protease was loaded in the well followed by 24 hours of incubation at room temperature.

Application of Proteases in detergents

Commercial Detergent Compatibility of the Partially Purified Protease

Using a concentration of 1%, different commercial detergents such as Ariel, Surf Excel, Tide and Rin were used to test the enzyme's stability in the presence of detergents. First, the enzymes that had been present in the detergent were heat-inactivated by boiling it for ten minutes. Skim milk agar wells were filled with the commercial detergents and the test enzyme solution.

Studies on Wash Performance

On a piece of white cotton fabric stained with dal and blood, the wash performance of the partly purified protease was investigated. After being stained with dal and human blood, the textile piece was allowed to dry for 12 hours. The pieces of stained cloth were placed on different Petri plates and exposed to the wash treatment experiments. The cloth pieces were removed from the incubator after an hour at room temperature, washed under tap water, allowed to dry, and then visually inspected to determine whether the stain had been effectively removed.

RESULTS AND DISCUSSION

Protease Production Screening

The positive strains were evaluated for protease production using 2 % skim milk agar. All positive strains were inoculated separately on skim milk agar to screen one bacterium with higher protease production. The single strain with a wider zone of clearance was used for further studies (Figure: 3.1).

Agarose Gel Electrophoresis of Isolated DNA and PCR Amplification

The DNA marker and the extracted DNA from the positive strain were seen on an agarose gel. A good DNA yield was shown by the brilliant appearance of the isolated DNA from the sample on lane 2 of the gel (Fig: 3.2).

Multiple sequence alignment of Isolated DNA

The BLAST results indicated that the obtained 16S rRNA gene sequence is matched with the maximum sequence homology with bacillaceae family. Phylogenetic trees were constructed by neighbor joining method. The tested strain showed maximum similarity with *Bacillus subtilis subsp. subtilis* strain KS6-27 and (Figure: 3.3) represents the phylogenetic tree of identified positive strain *Bacillus subtilis* AD20 constructed using mega 6.0 software with available strains in GenBank data (NCBI). The genbank accession number of the positive isolate is MT322145.

Augmentation of Culture Conditions for maximum Protease Production Influence of pH on Protein Content and Enzyme Activity

The results showed that the pH significantly influenced the protein content and protease activity in *Bacillus subtilis* AD20. It was able to relate the maximum protein content of 146 μ g/ml at pH 7 on 3rd day (Figure: 3.4.1a). At the same conditions, protease activity of 20,204.08 U/mg was obtained (Figure: 3.4.1b). The extracellular protein and





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protease activity production at different pH revealed that at pH 7 protease activity was highly significant than other pH. Therefore, pH 7 was selected for further studies. This indicated that the specific strain was alkaliphilic in nature and optimum pH 7 for growth and enzyme production is an ideal character among alkaliphilic organisms [11, 12]. It is commonly recognized that the permeability of bacterial cell membranes and the availability of specific metabolic ions are influenced by the pH of the culture medium, and that these factors promote the development of cells and the synthesis of enzymes [13].

Effect of Different Time Duration on Protein Content and Protease Activity

The *Bacillus subtilis* AD20 produced a maximum protein content of $90.5 \mu g/ml$ on the 3rd day (Figure: 3.4.2a) and protease activity of 30,400 U/mg on the 3rd day at 55° C (Figure: 3.4.2b) of incubation. The protein content and protease activity showed that 55° C temperature on the 3rd day was highly significant than other days. Thus 3rd day was selected for further studies. As the incubation period extended, both cell growth and enzyme synthesis rose linearly, indicating that growth and enzyme production are naturally correlated [12, 14]. As previously reported, after 72 hours of incubation, the best growth and enzyme production were seen [15, 16].

Impact of Various Carbon Sources on Protein and Protease Activity

When dextrose was tested at 1 % for protease production in *Bacillus subtilis* AD20, it showed an enhanced protein content of 120 μ g/ml on 3rd day and it induced maximum protease activity of 40,000 U/mg on 3rd day. Whereas minimum enzyme activity of 11,858.40 U/mg on 4th day in fructose and was not much significantly induced protease production. The protein content and protease activity production in different carbon sources revealed that dextrose was found highly significant than other carbon sources.

Impact of Various N2 Sources on Protein content and Protease Activity

Five different 1 % nitrogen sources such as ammonium acetate, yeast extract, peptone, casein and urea were taken. Nitrogen source when tested at 1 % for enzyme production in Bacillus subtilis AD20 it showed a maximum protein content of 129 µg/ml on 3rd day with ammonium acetate. But with casein, higher protein content of 124 µg/ml on 3rd day and optimum protease activity of 23,621.62 U/mg was observed. Whereas, minimum protease activity of 4,496.12 U/mg on 1st day in urea and 5,368.42 U/mg in yeast extract recorded on the 4th day was observed. The protein content and protease production in different nitrogen sources revealed that casein was found highly significant than other nitrogen sources. Since nitrogen is the final building block of protein biosynthesis, its availability has a significant impact on the synthesis of enzymes. Additionally, the medium's pH can be impacted by the nitrogen source, which could have an impact on the enzyme's stability and activity [17]. The similar outcome was earlier reported by Gajju et al. (1996), who discovered that because organic nitrogen sources are rich in amino acids, Bacillus sp. were more adapted to them for the manufacture of enzymes [18]. A few essential amino acids, some carbohydrates and inorganic substances like phosphorus and calcium are also present in casein.

Impact of Various substrate on Protein and Protease Activity

Five different substrates such as 1 % green gram husk, rice bran, sugarcane bagasse, grape waste, and kitchen waste was taken. Rice straw, when tested at 1 % for protease production in Bacillus subtilis AD20 showed high protein content of 82 μ g/ml on the 4th day and it induced maximum protease activity of 21,754.38 U/ml on 3rd day. The maximum protein content of 174 μ g/ml on 3rd day in green gram husk was observed. Whereas minimum enzyme activity of 6,542.05 U/ml on 1st day in kitchen waste and 7,032.25 U/ml in grape waste recorded on the 4th day of incubation was seen. The protein content and protease production in different substrates revealed that rice straw was found highly significant than other carbon sources.

Protease Production in Laboratory Scale

In this work, ammonium sulfate precipitation (80% saturation) was used to partially purify the protease enzyme from *Bacillus subtilis* AD20. The amount of extracellular protease production by *Bacillus sp.* was estimated on a laboratory scale from optimized parameters. The protease activity reached a maximum of 790 U/ml on the 3rd day of incubation. The crude culture fitrate containing enzyme was exposed to the precipitation method like ammonium





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sulfate to precipitate, partially purify and concentrate total protein. Finally, the precipitated enzyme was dialyzed and used for further analysis. The resulting partially purified protease had a specific activity of 24,687.5 U/mg. The molecular weight of protease was reported to be 35 kDa while the protease in literature was found to be 47 kDa [19]. The molecular mass of alkaline serine proteases normally comes in the array of 18 to 30 kDa [20] with few exceptions.

Detection of Extracellular Protease Profile on SDS-PAGE

SDS-PAGE was used to analyze the samples obtained at the partially purified stage. Following an electrophoretic run, 15% of the gel was stained with SDS-polyacrylamide using G-250 Coomassie staining (Figure: 3.5). Because complex medium has a specific substrate under growth conditions, only two bands were seen on the gel. These two bands on the gel, which ranged in size from 40 kDa to 30 kDa, may be used to calculate the molecular weight of the enzymes.

Detection of Extracellular Protease Profile on Native-PAGE

On a 1% gelatin gel, the protein band that was corresponding for enzymatic activity demonstrated creating the halozone (Figure: 3.6). This process was performed after the ammonium sulfate precipitated protease was separated on Native-PAGE.

The protein marker bands are indicated by lane-2 arrows, while lane-1 indicates the clearance of protease bands (Figure: 3.6). Along the gel's side is a line that indicates each protein marker band's molecular weight. According on Native-PAGE analysis, the isolated proteases had a molecular weight of about 30 kDa. On the gelatin gel, there was just one clearing zone that could be seen, signifying a single kind of protease. A gel was cleared by partially purified protease in accordance with the protein marker's electrophoretic mobility. Proteases from bacteria typically have a molecular weight of 15–36 kDa [21]. The molecular weight of B. mojavensis protease was 30 kDa, greater than that of alkaline protease [22]. Both proteases, however, are not as large as the Bacillus sp. strain GX6638's serine version of enzyme HS (36 kDa) [23].

Physicochemical Properties of Partially Purified Protease pH Optimum and Stability of Partially Purified Protease

The optimal pH for the activity of protease was pH 4 i.e., acidic pH. As pH goes on increasing, the relative activity of the enzyme goes on decreasing (Figure: 3.7). This optimal pH value is similar to the endoglucanase from Bacillus sp. [24]. The efficient operation of the enzyme was impacted as the pH value strayed from the ideal range. This might be attributed to either a lower affinity or reduced enzyme saturation with the substrate, or to the impact of pH on the stability of the enzyme [25].

Temperature Optimum and Stability

Higher temperatures are necessary for enzymes used in industries to remain active and stable, and temperature plays a vital role in maximizing enzyme activity. The partially purified protease of *Bacillus subtilis* AD20 showed a maximum enzyme activity at 55° C to determine its thermal stability, the enzyme was pre-incubated from 4° C to 55° C for 1 hour. It was stable from 24° C to 37° C with more 85.92 % to 90.37 % activity up to 60 minutes of incubation, respectively. Whereas, the protease at 55° C showed 100 % activity up to 1 hour (Figure: 3.8). On the other hand, the temperature optima of bacterial proteases are greater, reaching up to 85°C [26] and 115°C [27]. According to Bidochka and Khachatourians (1987), proteases function best at temperatures between 37 and 45°C [28].

Substrate Specificity of Partially Purified Protease

Among the specific substrates, casein was suitable for protease production. Peptone also showed good protease production when compared to others, but less then casein. The least protease production with the gelatin substrate (Figure: 3.9). It has been shown that the proteases from *Beauveria bassiana* and *Conidiobolus* sp. (*Cephalosporium* sp. KM388) are more active against casein than they are against bovine serum albumin. The catalytic process of an enzyme is modified by the presence of organic solvents due to the breaking of hydrogen bonds and hydrophobic interactions, as well as modifications in the protein's kinetics and structure.





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Organic Solvent of Partially Purified Protease

Among all organic solvents, benzene was appropriate for enzyme production when compared to other organic solvents i.e., ethanol, methanol, and DMSO. The least favorable for protease production was methanol and DMSO (Figure: 3.10). The physicochemical properties of partially purified protease were optimized with pH 4 i.e., acidic pH, 55° C determines its thermal stability, casein as substrates, and organic solvent benzene with maximum relative activity among other parameters. According to Gupta and Khare (2007), Pseudomonas aeruginosa protease was found to be relatively less stable in the presence of a hydrophilic solvent and more stable in organic solvents such as benzene, isooctane, cyclohexane and n-dodecane [29]. We can therefore assume that the type of organic solvents used will determine how stable the enzyme is in organic solvents. This indicates that an enzyme's structure can occasionally be stabilized by substituting some of its water molecules with organic ones.

Partially Purified Enzyme Activity on Skim Milk with Different Concentration

The skim milk activity with different concentrations of partially purified protease was loaded and after incubation, the zone of clearance was observed. The 30 μ l concentration showed a higher zone of clearance when compared with different other concentrations. The least zone was observed in 20 μ l of concentration of partially purified protease (Figure: 3.11). The caseinolytic activity was also performed but no results were detected in the casein agar plates.

Application of Partially Purified Protease Wash Performance Test

The comparison of washing of stained clothes with dal and blood after 12 hours of drying the stain was done. These clothes after incubation with distilled water and partially purified protease were observed. The clothes in partially purified protease were not having the outer stains and stains of both blood and dal was removed more than the distilled water (Figure: 3.12). Hence, the results showed that the enzyme can be used commercially for detergent purposes.

Comparison of Partially Purified Protease and Commercial Detergents

The activity of commercial detergent and partially purified protease was performed in skim milk agar. This showed the test sample was comparatively positive when compared to the detergent activity by the zone of clearance. This zone of clearance was higher in Ariel than the test sample, with the decreasing activity with the other detergents in the similar concentrations (Figure: 3.13). Protease from *Conidiobolus coronatus* was found to be stable and compatible when added to commercial detergents, according to Phadtare et al. (1993). There is an increasing need to investigate bacterial proteases for commercially viable applications in the detergent industry, as there are limited reports on the use of fungal proteases in this sector [30]. Protease from *Virgibacillus pantothenticus* that is stable in the presence of readily available detergents locally was reported by Gupta et al. (2008). Three hours of incubation at 40°C produced more than 50% activity in the protease [31].

CONCLUSION

In conclusion, the investigations resulted in a partial purification and characterization of the protease isolated from *Bacillus subtilis* AD20. The bacterially generated enzyme has an acidic and rather thermostable composition. In further research, chromatography can be used to purify it, and it can be immobilized on either organic or inorganic substrate for use in commercial operations. To increase its use in biotechnology, it can also be made by recombinant DNA technology. Future research should focus on X-ray crystallographic examinations of the proteases, since these will provide insight into the characteristics of the catalytic and structural metal-binding sites.





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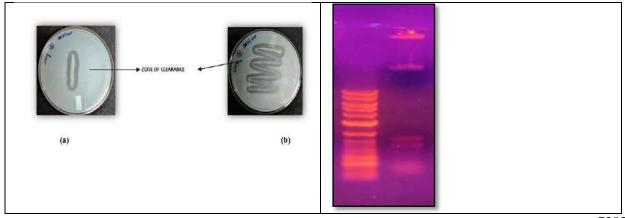




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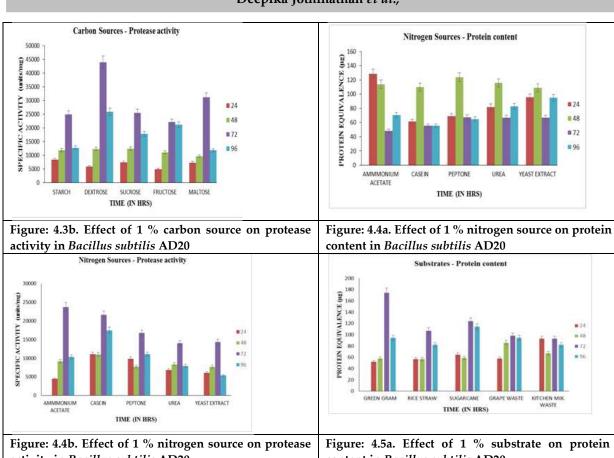
Figure 1. Screening of Single Positive Strain on 2 Figure 2.a. Agarose Gel Electrophoresis of PCR %Skim Milk Agar in (a) and (b) product MT184827.1 Bacillus subtilis strain NWPZ-15 MN044860.1 Bacillus velezensis strain ANKO3 pH - Protein content MK736123.1 Bacillus sp. (in: Bacteria) strain SS1 MN444025.1 Bacillus subtilis strain A4 200 MK564708.1 Bacillus subtilis strain DCH4 MK729017.1 Bacillus tequilensis strain J731 EQUIVALENCE (µg) 150 MN893860.1 Bacillus vallismortis strain HR16 **24** MH187568.1 Bacillus sp. strain LLB-07 100 MN067800.1 Bacillus subtilis strain PaKu23 **PROTEIN** MN907664.1 Bacillus tequilensis strain SBRSW24 **48** 50 MN493056.1 Bacillus subtilis strain NG106 MN889298.1 Bacillus subtilis strain OsEnb_PLM_S6 0 **72** MT211515.1 Bacillus tequilensis strain SKC-10 MN905164.1 Bacillus subtilis strain Bs 2 3 5 1 -MN946550.1 Bacillus subtilis subsp. subtilis strain KS6-27 **96** MT322145 Bacillus subtilis AD20 TIME (IN HRS) Figure 3. Phylogenetic position of Bacillus subtilis Figure: 4.1 a. Effect of different pH on protein AD20 using neighbors-joining method with the content in Bacillus subtilis AD20 related genus pH - Protease activity **TIME - PROTEIN CONTENT** 100 30000 EQUIVALENCE (μg) PECIFIC ACTIVITY 80 20000 **2**4 60 10000 (units/mg) **48** 40 **7**2 20 0 96 TIME (IN HRS) 24 48 72 96 TIME (IN HOURS) Figure: 4.1 b. Effect of different pH on protease Figure: 4.2a. Effect of time duration on protein activity in Bacillus subtilis AD20 content in Bacillus subtilis AD20 Carbon Sources - Protein content **TIME - PROTEASE** 140 ACTIVITY B 120 40,000.00 PROTEIN EQUIVALENCE 100 SPECIFIC ACTIVITY 30,000.00 20,000.00 24hrs (units/mg) 10,000.00 0.00 m 72hrs 24 48 **72** 96 TIME (IN HOURS) DEXTROSE SUCROSE FRUCTOSE TIME (IN HRS) Figure: 4.2b. Effect of time duration on protease Figure: 4.3a. Effect of 1 % carbon source on protein activity in Bacillus subtilis AD20 content in Bacillus subtilis AD20





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activity in Bacillus subtilis AD20

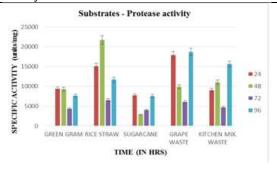


Figure: 4.5b. Effect of 1 % substrate on protease activity in Bacillus subtilis AD20

Figure: 4.5a. Effect of 1 % substrate on protein content in Bacillus subtilis AD20

PROTEIN MARKER CRUDE DIALYSED SAMPLE

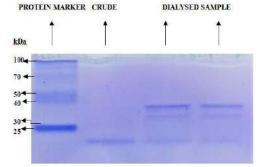
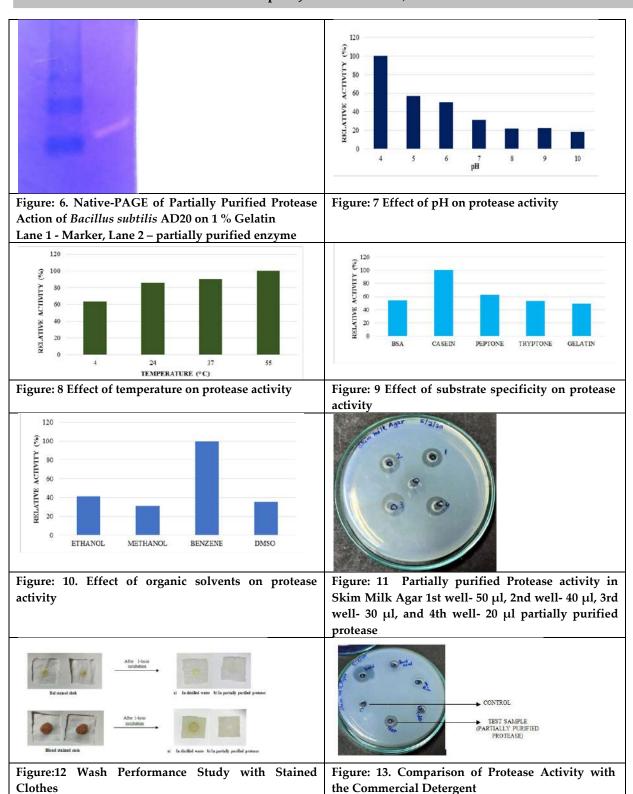


Figure: 5. SDS-PAGE with Coomassie Staining G-250





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RESEARCH ARTICLE

Homomorphism in Bipolar Valued Vague Subnear Rings of a Nearring

S. Kamalaselvi¹ and B.Anandh^{2*}

¹Research Scholar, Department of Mathematics, Sudharsan College of Arts and Science (Affiliated to Bharathidasan University, Tiruchirappalli), Pudukkottai, Tamil Nadu, India.

²Assistant Professor, Department Mathematics, H. H. The Rajah's College (Affiliated to Bharathidasan University, Tiruchirappalli), Pudukkottai, Tamil Nadu, India.

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*Address for Correspondence

B.Anandh

Assistant Professor, Department Mathematics,

H. H. The Rajah's College

(Affiliated to Bharathidasan University, Tiruchirappalli),

Pudukkottai, Tamil Nadu, India. Email: drbaalaanandh@gmail.com



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ABSTRACT

In this paper, bipolar valued vague subnearring of a nearring is studied by homomorphism and anti homomorphism and some properties are discussed. These properties are useful to further research.

Keywords: Fuzzy subset, vague subset, bipolar valued fuzzy subset, bipolar valued vague subset, bipolar valued vague subnearring, bipolar valued vague normal subnearring, intersection, image and preimage.

INTRODUCTION

In 1965, Zadeh [13] introduced the notion of a fuzzy subset of a Universal set. Vague set is an extension of fuzzy set and it is appeared as a unique case of context dependent fuzzy sets. The vague set was introduced by W.L.Gau and D.J.Buehrer [5]. W.R.Zhang [14, 15] introduced an extension of fuzzy sets named bipolar valued fuzzy sets in 1994 and bipolar valued fuzzy set was developed by Lee [6, 7]. Fuzzy subgroup was introduced by Azriel Rosenfeld [3]. RanjitBiswas [9] introduced the vague groups. Cicily Flora. S and Arockiarani. I [4] have introduced a new class of generalized bipolar vague sets. Anitha.M.S., et.al.[1] defined as bipolar valued fuzzy subgroups of a group. Sheena. K. P and K.Uma Devi [10] have introduced the bipolar valued fuzzy subbigroup of a bigroup. Shanthi.V.K and G.Shyamala[11] have introduced the bipolar valued multi fuzzy subgroups of a group. Yasodara.S, KE. Sathappan [12] defined the bipolar valued multi fuzzy subsemirings of a semiring. Bipolar valued multi fuzzy subnearring of a nearing has been introduced by S.Muthukumaran and B.Anandh [8]. Anitha.K., et.al.[2] defined as bipolar valued vague subsemirings of a semiring. Here, the concept of bipolar valued vague subnearring of a nearring is introduced





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and estaiblished some results. Homomorphism and anti homomorphism are applied in bipolar valued vague subnearring of a nearring.

PRELIMINARIES

Definition 1.1. [13] Let X be any nonempty set. A mapping M : $X \to [0, 1]$ is called a fuzzy subset of X. **Definition 1.2. [5]** A vague set A in the universe of discourse U is a pair $[t_A, 1-f_A]$, where $t_A : U \to [0, 1]$ and $f_A : U \to [0, 1]$ are mappings, they are called truth membership function and false membership function respectively. Here $t_A(x)$ is a lower bound of the grade of membership of x derived from the evidence for x and $f_A(x)$ is a lower bound on the negation of x derived from the evidence against x and $t_A(x) + f_A(x) \le 1$, for all $x \in U$.

Definition 1.3. [5] The interval [$t_A(x)$, $1-f_A(x)$] is called the vague value of x in A and it is denoted by $V_A(x)$, i.e., $V_A(x) = [t_A(x), 1-f_A(x)]$.

Example 1.4. A = $\{< a, [0.1, 0.2] >, < b, [0.3, 0.4] >, < c, [0.5, 0.6] >\}$ is a vague subset of $X = \{a, b, c\}$.

Definition 1.5. [14] A bipolar valued fuzzy set (BVFS) A in X is defined as an object of the form $A = \{ < x, A^+(x), A^-(x) > / x \in X \}$, where $A^+ : X \to [0, 1]$ and $A^- : X \to [-1, 0]$. The positive membership degree $A^+(x)$ denotes the satisfaction degree of an element x to the property corresponding to a bipolar valued fuzzy set A and the negative membership degree $A^-(x)$ denotes the satisfaction degree of an element x to some implicit counter-property corresponding to a bipolar valued fuzzy set A.

Example 1.6. A = {<a, 0.2, -0.3>, <b, 0.4, -0.5>, <c, 0.6, -0.7>} is a bipolar valued fuzzy subset of X = {a, b, c}. **Definition 1.7. [4]** A bipolar valued vague subset A in X is defined as an object of the form A = { $\langle x, [t_A^+(x), 1-f_A^+(x)], [-1-f_A^-(x), t_A^-(x)] \rangle / x \in X$ }, where $t_A^+: X \to [0, 1]$, $f_A^+: X \to [0, 1]$, $t_A^-: X \to [-1, 0]$ and $f_A^-: X \to [-1, 0]$ are mapping such that $t_A(x) + f_A(x) \le 1$ and $-1 \le t_A^- + f_A^-$. The positive interval membership degree [$t_A^+(x), 1-f_A^+(x)$] denotes the satisfaction region of an element x to the property corresponding to a bipolar valued vague subset A and the negative interval membership degree [$-1-f_A^-(x), t_A^-(x)$] denotes the satisfaction region of an element x to some implicit counter-property corresponding to a bipolar valued vague subset A. Bipolar valued vague subset A is denoted as A = { $\langle x, V_A^+(x), V_A^-(x) \rangle / x \in X$ }, where $V_A^+(x), = [t_A^+(x), 1-f_A^+(x)]$ and $V_A^-(x) = [-1-f_A^-(x), t_A^-(x)]$.

Note that. [0] = [0, 0], [1] = [1, 1]and [-1] = [-1, -1].

Example 1.8. [A] = { < a, [0.5, 0.8], [-0.4, -0.1] >, < b, [0.22, 0.54], [-0.7, -0.2] >, < c, [0.11, 0.5], [-0.8, -0.5] >} is a bipolar valued vague subset of X = {a, b, c}.

Definition 1.9. [4] Let $A = \langle V_A^+, V_A^- \rangle$ and $B = \langle V_B^+, V_B^- \rangle$ be two bipolar valued vague subsets of a set X. We define the following relations and operations:

- (i) $A \subset B$ if and only if $V_A^+(u) \le V_B^+(u)$ and $V_A^-(u) \ge V_B^-(u)$, $\forall u \in X$.
- (ii) A = B if and only if V_A^+ (u) = V_B^+ (u) and V_A^- (u) = V_B^- (u), \forall u \in X.
- (iii) $A \cap B = \{ \langle u, rmin(V_A^+(u), V_B^+(u)), rmax(V_A^-(u), V_B^-(u)) \rangle / u \in X \}.$





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 $\begin{aligned} & \text{(iv) } A \cup B = \{ \ \langle \ \mathbf{u}, \ \text{rmax } (V_A^+(\mathbf{u}), \ V_B^+(\mathbf{u}) \), \ \text{rmin } (V_A^-(\mathbf{u}), \ V_B^-(\mathbf{u}) \) \ \rangle \ / \ \mathbf{u} \in \mathbf{X} \ \}. \ \text{Here rmin } (V_A^+(\mathbf{u}), \ V_B^+(\mathbf{u}) \) = [\ \min \{ t_A^+(x), t_B^+(x) \ \}, \ \min \{ 1 - f_A^+(x), 1 - f_B^+(x) \} \], \ \text{rmax } (V_A^+(\mathbf{u}), \ V_B^+(\mathbf{u}) \) = [\ \max \{ t_A^+(x), t_B^+(x) \ \}, \ \max \{ 1 - f_A^+(x), t_B^-(x) \} \], \ \text{rmax } (V_A^-(\mathbf{u}), \ V_B^-(\mathbf{u}) \) = [\ \max \{ -1 - f_A^-(x), -1 - f_B^-(x) \}, \ \max \{ t_A^-(x), t_B^-(x) \} \]. \end{aligned}$

Definition 1.10. Let R be a nearring. A bipolar valued vague subset A of R is said to be a bipolar valued vague subnearring of R (BVVSNR) if the following conditions are satisfied,

- (i) $V_A^+(x-y) \ge \text{rmin}\{V_A^+(x), V_A^+(y)\}$
- (ii) $V_A^+(xy) \ge \text{rmin}\{ V_A^+(x), V_A^+(y) \}$
- (iii) $V_A^-(x-y) \le \text{rmax}\{ V_A^-(x), V_A^-(y) \}$
- (iv) $V_A^-(xy) \le \text{rmax}\{ V_A^-(x), V_A^-(y) \}$ for all x and y in R.

Example 1.11. Let $R = Z_3 = \{0, 1, 2\}$ be a nearring with respect to the ordinary addition and multiplication. Then $A = \{0, [0.6, 0.8], [-0.9, -0.6] >, <1, [0.5, 0.7], [-0.8, -0.5] >, <2, [0.5, 0.7], [-0.8, -0.5] >\}$ is a BVVSNR of R.

Definition 1.12. Let R be a nearring. A BVVSNR A = $\langle V_A^+, V_A^- \rangle$ of R is said to be a bipolar valued vague normal subnearring (BVVNSNR) of R if V_A^+ (x+y) = V_A^+ (y+x), V_A^+ (xy) = V_A^+ (yx), V_A^- (x+y) = V_A^- (y+x) and V_A^- (xy) = V_A^- (yx) for all x and y in R.

Definition 1.13. Let R and R¹ be any two nearrings. Then the function $f: R \to R^1$ is said to be an antihomomorphism if f(x+y) = f(y) + f(x) and f(xy) = f(y) + f(x) for all x and y in R.

Definition 1.14. Let X and X^I be any two sets. Let $f: X \to X^I$ be any function and let $A = \langle V_A^+, V_A^- \rangle$ be a bipolar valued vague subset in X, $V = \langle V_V^+, V_V^- \rangle$ be a bipolar valued vague subset in $f(X) = X^I$, defined by $V_V^+(y) = r \inf_{x \in f^{-1}(y)} V_A^-(x)$, for all x in X and y in X^I. A is called a preimage of V under f and is

defined as $V_A^+(x) = V_V^+(f(x))$, $V_A^-(x) = V_V^-(f(x))$ for all x in X and is denoted by f⁻¹(V).

SOME THEOREMS.

Theorem 2.1. Let R and R' be any two nearrings. The homomorphic image of a BVVSNR of R is a BVVSNR of R'.

Proof: Let f : R→ R¹ be a homomorphism. Let V = f(A) = $\langle V_V^+, V_V^- \rangle$, where A = $\langle V_A^+, V_A^- \rangle$ is a BVVSNR of R. We have to prove that V is a BVVSNR of R¹. Now for f(x), f(y) in R¹, V_V^+ (f(x)-f(y)) = V_V^+ (f(x-y)) ≥ V_A^+ (x-y) ≥ rmin{ V_A^+ (x), V_A^+ (y)} = rmin{ V_V^+ (f(x)), V_V^+ (f(y)) } which implies that V_V^+ (f(x)-f(y)) ≥ rmin{ V_V^+ (f(x)), V_V^+ (f(y))}. And V_V^+ (f(x)f(y)) = V_V^+ (f(xy)) ≥ V_A^+ (xy) ≥ rmin{ V_A^+ (x), V_A^+ (y)} = rmin { V_V^+ (f(x)), V_V^+ (f(y)) } which implies that V_V^+ (f(x)f(y)) ≥ rmin{ V_V^+ (f(x)), V_V^+ (f(y)) }. Also V_V^- (f(x)-f(y)) = V_V^- (f(x-y)) ≤ V_A^- (x-y) ≤ rmax { V_A^- (x), V_A^- (y)} = rmax { V_V^- (f(x)), V_V^- (f(y))} which implies that V_V^- (f(x)-f(y)) ≤ rmax{ V_V^- (f(x)), V_V^- (f(y)) }. And V_V^- (f(x)f(y)) = V_V^- (f(x)). Which implies that V_V^- (f(x)f(y)) ≤ rmax{ V_V^- (f(x)), V_V^- (f(y)) }. Hence V is a BVVSNR of R¹.

Theorem. Let R and R † be any two nearrings. The homomorphic preimage of a BVVSNR of R † is a BVVSNR of R.





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Proof. Let f : R → R¹ be a homomorphism. Let V = f(A) = $\langle V_V^+, V_V^- \rangle$ where V is a BVVSNR of R¹. We have to prove that A = $\langle V_A^+, V_A^- \rangle$ is a BVVSNR of R. Let x and y in R. Now V_A^+ (x-y) = V_V^+ (f(x-y)) = V_V^+ (f(x)-f(y)) ≥ rmin { V_V^+ (f(x)), V_V^+ (f(y)) }= rmin { V_A^+ (x), V_A^+ (y)} which implies that V_A^+ (x-y) ≥ rmin{ V_A^+ (x), V_A^+ (y) }. And V_A^+ (xy) = V_V^+ (f(xy)) = V_V^+ (f(x)f(y)) ≥ rmin{ V_V^+ (f(x)), V_V^+ (f(y)) }= rmin { V_A^+ (x), V_A^+ (y)} which implies that V_A^+ (xy) ≥ rmin{ V_A^+ (x), V_A^+ (y) }. Also V_A^- (x-y) = V_V^- (f(x-y)) = V_V^- (f(x)-f(y)) ≤ rmax{ V_V^- (f(x)), V_V^- (f(y))} = rmax{ V_A^- (x), V_A^- (y)} which implies that V_A^- (x-y) ≤ rmax{ V_A^- (x), V_A^- (y) }. And V_A^- (xy) = V_V^- (f(xy)) = V_V^- (f(x)f(y)) ≤ rmax{ V_V^- (f(x)), V_V^- (f(y))} = rmax{ V_A^- (x), V_A^- (y)} which implies that V_A^- (x), V_A^- (y)} which implies that V_A^- (x), V_A^- (y)}. Hence A is a BVVSNR of R.

Theorem: Let R and R¹ be any two nearrings. The antihomomorphic image of a BVVSNR of R is a BVVSNR of R¹.
Proof: Let f: R → R¹ be an antihomomorphism. Let V = f(A) = $\langle V_V^+, V_V^- \rangle$ where $A = \langle V_A^+, V_A^- \rangle$ is a BVVSNR of R.
We have to prove that V is a BVVSNR of R¹. Now for f(x), f(y) in R¹, V_V^+ (f(x) −f(y)) = V_V^+ (f(y−x)) ≥ V_A^+ (y−x) ≥ rmin { V_A^+ (x), V_A^+ (y) } = rmin { V_V^+ (f(x)), V_V^+ (f(y)) } which implies that V_V^+ (f(x) −f(y)) ≥ rmin{ V_V^+ (f(x)), V_V^+ (f(y)) }. And V_V^+ (f(x)f(y)) = V_V^+ (f(x)), V_V^+ (f(y)) }. Also V_V^- (f(x) −f(y)) = V_V^- (f(x)), V_V^+ (f(y)) } which implies that V_V^+ (f(x)f(y)) ≥ rmin { V_V^+ (f(x)), V_V^+ (f(y)) }. Also V_V^- (f(x) −f(y)) = V_V^- (f(x)), V_V^- (f(y))}. And V_V^- (f(x)f(y)) = rmax{ V_V^- (f(x)), V_V^- (f(y))} which implies that V_V^- (f(x) −f(y)) ≤ rmax{ V_V^- (f(x)), V_V^- (f(y))}. And V_V^- (f(x)f(y)) ∈ V_V^- (f(x)f(y)) } = V_V^- (f(x)), V_V^- (f(x)). Hence V is a BVVSNR of R¹.

Theorem: Let R and R¹ be any two nearrings. The antihomomorphic preimage of a BVVSNR of R¹ is a BVVSNR of R. Proof: Let f: R → R¹ be an antihomomorphism. Let V = f(A) = $\langle V_V^+, V_V^- \rangle$ where V is a BVVSNR of R¹. We have to prove that A = $\langle V_A^+, V_A^- \rangle$ is a BVVSNR of R. Let x and y in R. Now V_A^+ (x-y) = V_V^+ (f(x-y)) = V_V^+ (f(y) -f(x)) ≥ rmin{ V_V^+ (f(x)), V_V^+ (f(y)) } = rmin{ V_A^+ (x), V_A^+ (y) } which implies that V_A^+ (x-y) ≥ rmin{ V_A^+ (x), V_A^+ (y)}. And V_A^+ (xy) ≥ rmin{ V_A^+ (x), V_A^+ (y)}. Also V_A^- (x-y) = V_V^- (f(x-y)) = V_V^- (f(y) -f(x)) ≤ rmax{ V_V^- (f(x)), V_V^- (f(y)) } = rmax{ V_A^- (x), V_A^- (y)} which implies that V_A^- (x), V_A^- (y)}. And V_A^- (x) = rmax{ V_A^- (x), V_A^- (y)} which implies that V_A^- (x), V_A^- (y)}. Hence A is a BVVSNR of R.

Theorem: Let R and R¹ be any two nearrings. The homomorphic image of a BVVNSNR of R is a BVVNSNR of R¹. **Proof.** Let f: R \rightarrow R¹ be a homomorphism. Let V = f(A) = $\langle V_V^+, V_V^- \rangle$ where A = $\langle V_A^+, V_A^- \rangle$ is a BVVNSNR of R. We have to prove that V is a BVVNSNR of R¹. By Theorem 2.1, V is a BVVSNR of R¹. Now for f(x), f(y) in R¹, V_V^+ (f(x)+f(y)) = V_V^+ (f(x+y)) $\geq V_A^+$ (x+y) = V_A^+ (y+x) $\leq V_V^+$ (f(y+x)) = V_V^+ (f(y)+f(x)) which implies that V_V^+ (f(x)+f(y)) =





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 $V_{V}^{+}\left(\mathbf{f}(\mathbf{y})+\mathbf{f}(\mathbf{x})\right). \text{ And } V_{V}^{+}\left(\mathbf{f}(\mathbf{x})\mathbf{f}(\mathbf{y})\right) = V_{V}^{+}\left(\mathbf{f}(\mathbf{x}\mathbf{y})\right) \geq V_{A}^{+}\left(\mathbf{x}\mathbf{y}\right) = V_{A}^{+}\left(\mathbf{y}\mathbf{x}\right) \leq V_{V}^{+}\left(\mathbf{f}(\mathbf{y}\mathbf{x})\right) = V_{V}^{+}\left(\mathbf{f}(\mathbf{y})\mathbf{f}(\mathbf{x})\right) \text{ which implies that } V_{V}^{+}\left(\mathbf{f}(\mathbf{x})\mathbf{f}(\mathbf{y})\right) = V_{V}^{+}\left(\mathbf{f}(\mathbf{y})\mathbf{f}(\mathbf{x})\right). \text{ Also } V_{V}^{-}\left(\mathbf{f}(\mathbf{x})+\mathbf{f}(\mathbf{y})\right) = V_{V}^{-}\left(\mathbf{f}(\mathbf{x}+\mathbf{y})\right) \geq V_{A}^{-}\left(\mathbf{x}+\mathbf{y}\right) = V_{A}^{-}\left(\mathbf{y}+\mathbf{x}\right) \leq V_{V}^{-}\left(\mathbf{f}(\mathbf{y})+\mathbf{f}(\mathbf{x})\right) = V_{V}^{-}\left(\mathbf{f}(\mathbf{y})+\mathbf{f}(\mathbf{y})\right) = V_{V}^{$

Theorem: Let R and R¹ be any two nearrings. The homomorphic preimage of a BVVNSNR of R¹ is a BVVNSNR of R. Proof: Let f: R \rightarrow R¹ be a homomorphism. Let V = f(A) = $\langle V_V^+, V_V^- \rangle$ where V is a BVVNSNR of R¹. We have to prove that A = $\langle V_A^+, V_A^- \rangle$ is a BVVNSNR of R. By Theorem 2.2, A = $\langle V_A^+, V_A^- \rangle$ is a BVVSNR of R. Let x and y in R. Now V_A^+ (x+y) = V_V^+ (f(x+y)) = V_V^+ (f(x)+f(y)) = V_V^+ (f(y)+f(x)) = V_V^+ (f(y+x)) = V_A^+ (y+x) which implies that V_A^+ (x+y) = V_A^+ (y+x). And V_A^+ (xy) = V_V^+ (f(xy)) = V_V^+ (f(x)f(y)) = V_V^+ (f(y)f(x)) = V_V^+ (f(y)+f(x)) = V_V^- (f(y+x)) = V_A^- (yx) which implies that V_A^- (x+y) = V_A^- (y+x). And V_A^- (x+y) = V_V^- (f(x+y)) = V_V^- (f(x)+f(y)) = V_V^- (f(y)+f(x)) = V_V^- (f(y)f(x)) = V_V^- (f(x)f(x)) = $V_$

Theorem. Let R and R¹ be any two nearrings. The antihomomorphic image of a BVVNSNR of R is a BVVNSNR of R¹. **Proof:** Let f: R \rightarrow R¹ be an antihomomorphism. Let V = f(A) = $\langle V_V^+, V_V^- \rangle$ where A = $\langle V_A^+, V_A^- \rangle$ is a BVVNSNR of R.
We have to prove that V is a BVVNSNR of R¹. By Theorem 2.3, V is a BVVSNR of R¹. Now for f(x), f(y) in R¹, V_V^+ (f(x)+f(y)) = V_V^+ (f(y+x)) $\geq V_A^+$ (y+x) = V_A^+ (x+y) $\leq V_V^+$ (f(x+y)) = V_V^+ (f(y)+f(x)) which implies that V_V^+ (f(x)+f(y)) = V_V^+ (f(y)+f(x)). And V_V^+ (f(x)f(y)) = V_V^+ (f(yx)) $\geq V_A^+$ (yx) = V_A^+ (xy) $\leq V_V^+$ (f(xy)) = V_V^+ (f(y)f(x)) which implies that V_V^+ (f(x)f(y)) = V_V^+ (f(y)f(x)). Also V_V^- (f(x)+f(y)) = V_V^- (f(y+x)) $\leq V_A^-$ (y+x) = V_A^- (x+y) $\leq V_V^-$ (f(x+y)) = V_V^- (f(y)+f(x)) which implies that V_V^- (f(x)+f(y)) = V_V^- (f(x)f(y)). And V_V^- (f(x)f(y)) = V_V^- (f(xy)) $\leq V_A^-$ (yx) = V_V^- (f(xy)) which implies that V_V^- (f(x)) which imp

Theorem: Let R and R^{l} be any two nearrings. The antihomomorphic preimage of a BVVNSNR of R^{l} is a BVVNSNR of R.

Proof. Let $f: R \to R^1$ be an antihomomorphism. Let $V = f(A) = \langle V_V^+, V_V^- \rangle$ where V is a BVVNSNR of R^1 . We have to prove that $A = \langle V_A^+, V_A^- \rangle$ is a BVVNSNR of R. By Theorem 2.4, $A = \langle V_A^+, V_A^- \rangle$ is a BVVSNR of R. Let X and Y in R. Now $V_A^+(X+Y) = V_V^+(f(X+Y)) = V_V^+(f(Y)+f(X)) = V_V^+(f(X)+f(Y)) = V_V^+(f(Y+X)) = V_A^+(Y+X)$ which implies that $V_A^+(X+Y) = V_A^+(Y+X)$. And $V_A^+(X+Y) = V_V^+(Y+X) = V_V^-(Y+X) = V_V$





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RESEARCH ARTICLE

Ecotrail Energy Efficient Routing Algorithm (Eeera) for Green Internet of **Things**

A. Leolin Arockiadass^{1*}, T. Kokilavani², A. Aloysius³ and Y. Sunil Raj⁴

¹Research Scholar, Department of Computer Science, St. Joseph's College (Autonomous), (Affiliated to Bharathidasan University) Tiruchirappalli, Tamil Nadu, India.

²Assistant Professor, Department of Computer Science, CHRIST University, Bengaluru, Karnataka, India.

³Assistant Professor, Department of Computer Science, St. Joseph's College (Autonomous), (Affiliated to Bharathidasan University), Tiruchirappalli, Tamil Nadu, India.

⁴Assistant Professor, Department of Data Science, St. Xavier's College (Autonomous), Palayamkottai (Affiliated to Manonmaniam Sundaranar University), Tirunelveli, Tamil Nadu, India.

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*Address for Correspondence

A. Leolin Arockiadass

Research Scholar,

Department of Computer Science,

St. Joseph's College (Autonomous),

(Affiliated to Bharathidasan University)

Tiruchirappalli, Tamil Nadu, India.

Email: leolin.arul@gmail.com



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ABSTRACT

The advent of Green Internet of Things (G-IoT) heralds a transformative era in optimizing device utilization within smart environments, crucial for curtailing energy consumption and mitigating carbon emissions. This research addresses the pressing need for energy reduction, particularly at the network layer of the protocol stack, to foster eco-friendliness during data transmission. This work presents a novel routing algorithm, EcoTrial, leveraging Ant Colony Optimization (ACO) to bolster energy efficiency within G-IoT ecosystems. The simulation is conducted to evaluate EcoTrial's efficacy, revealing its superior performance compared to existing schemes. This pioneering approach not only advances sustainability goals but also diminishes environmental impact, thereby fostering the development of ecoconscious solutions in G-IoT environments.

Keywords: Green Internet of Things (G-IoT), Network Layer, Routing, Ant Colony Optimization (ACO), EcoTrial, Energy Efficiency





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INTRODUCTION

Smart devices with inherent capability to autonomously gather, process and exchange data among themselves and their ability to actuate a decision/ action in a given environment has put the Internet of Things on the frontier top of the technological map. The ubiquitous nature of IoT devices augments the immense possibilities for innovation and efficiency in various applications such as smart city, smart agriculture etc. The steady evolution of these devices has resulted in a new era of connectivity, automation, and data-driven decision-making[1]. From smart homes and cities to industrial processes and healthcare systems, the impact of IoT is far-reaching. The Indian subcontinent is witnessing fast urbanization and it is evident from the mushrooming of small, medium and very large sized cities with dense populations. As a consequence there is a steep upsurge in environmental degradation and energy consumption. There is also the need to provide urban services such as traffic congestion, sanitation and waste management. As the volume of IoT data and the amount of data transmitted is mounting up, the implementation of efficient and intelligent procedures that can lead to a reduction in energy consumption become imperative. [2] Routing techniques play a major role in the reduction of energy consumption in IoT implementations [3,4,5]. Routing holds the responsibility of choosing optimal communication paths and speed of transmission. Further its responsibility extends to preserving connection longevity in low-power and cost-effective scenarios [6, 7, 8]. Enhancing energy efficiency involves the application of optimization algorithms, with nature-inspired algorithms playing a predominant role. Techniques such as Ant Colony Optimization (ACO) are employed in conjunction with energy reduction strategies and scheduling techniques, addressing challenges such as the Traveling Salesman Problem [9, 10]. The work aims at proposing an algorithm that optimizes energy usage in IoT networks which lead towards the advancement of the G-IoT environment. The objectives include studying existing energy efficiency techniques based on ACO, and proposing a novel algorithm based on the ACO scheme. The remainder of this article follows a structured organization, the section 2, presents the Review of Literature, Section 3 delves into the proposed optimization technique coupled with ACO, Section 4, presents the results and analysis and finally section 5, draws conclusions.

REVIEW OF LITERATURE

Routing techniques in IoT networks play a major role be it heterogeneous or homogeneous networks. In heterogeneous networks, nodes are grouped into clusters, and the source node determines the intermediary node acting as a proxy across radio access technologies [11]. Optimization coupled with routing algorithms enhance energy consumption in IoT networks. Pushpender Saraol have evaluated and compared routing protocols such as AODV, DSR, and Destination-Sequenced Distance-Vector Routing (DSDV) [12]. ACO is prominent among optimization algorithms associated with routing protocols. The N-EEABR algorithm, an improvement over conventional Energy Efficient Ant-Based Routing, optimizes transmission power for enhanced efficiency in ant colony routing in IoT networks, promoting energy conservation and waste prevention [13]. Algorithms such as ACO-BFC-Minkowski Static (ABMS) and ACO-BFC-Minkowski Dynamic (ABMD), show significant energy savings and extend the network lifetime in static IoT scenarios. These two meta-heuristic approaches are optimized with Swarm Intelligence for shortest path computation in static and dynamic IoT networks. Utilizing widely-used algorithms like Breadth-First Search (BFS) and Dijkstra, the routes are further optimized using ACO[14]. Based on the study, the fact emerges that hybrid ACO enhances the energy efficiency in G-IoT while making the environment greener. As a result this work highly recommends optimization algorithms such as EcoTrial to be used in G-IoT architectures as it provides effective optimization, speed and efficiency in finding solutions over other recent algorithms.

Proposed Work

The proposed approach reduces the energy utilization of EcoTrial which combines Ant Colony Optimization (ACO) with Electric Fields in Nature (EFN). In EcoTrial, routing nodes, akin to ants, create virtual trails in a network influenced by an electric field. These nodes establish and maintain virtual trails to other nodes. Electric fields are





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formed between network nodes, where the source node carries a higher electric charge compared to the destination node. The strength of the electric field diminishes with increasing distance between nodes. Ant nodes prioritize trails representing the shortest path between source and destination, while also considering the influence of the electric field. They experience a force aligned with the electric field, influencing their routing decisions. As ant nodes traverse the network, they leave virtual "eco-trails" reflecting their chosen paths. The strength of these eco-trails is adjusted based on path quality factors such as energy consumption and latency. Furthermore, the electric field strength between nodes is updated based on traffic and energy consumption along the path.

The energy consumption of nodes during data transmission follows a linear model, as expressed by Equation 1, where E(s, d) denotes the energy consumed for transmitting from node s to node d, and is proportional to the distance D(s, d) between the nodes.

$$E(s, d) = \alpha \propto D(s, d)$$
 eq. 1

Equation 1, shows the energy consumed E(s, d) for transmitting a packet from node s to node d is proportional to the distance D(s, d) between the nodes. Here s is assumed to have higher mass and d will be with lower mass, α is a proportionality constant and α denotes proportionality. Figure 1, illustrates the connected components of an IoT network, depicting nodes as A, B, C, D, E, F, G, and H, with edges denoted by directed arrows and corresponding connections specified. The edges are connected with a directed arrows, where the connections can be represented as (A,B), (A,E), (B,C), (C,D), (D,E), (E,F), (F,G) and (G, H).

$$E(s, d) = \beta / (D(s, d))^2$$
 eq. 2

Equation 2 models the electric fields between nodes as inverse-square forces, where the electric field strength between nodes s and d is inversely proportional to the square of their distance, denoted by β .

Figure 2, illustrates the movement of packets, the edges marked with red signifies the max strength of pheromone (1), while the dashed red marking signifies the presence of pheromone at an average of 0.5. Subsequently the other edges without red marking signifies the unused routes whose pheromone count will be 0. The probability of an ant at node s moving to node d is determined by a combination of eco-trails and electric field strengths, as expressed in Equation 3, with the weight factor 'w' controlling the balance between exploration and exploitation. A higher 'w' emphasizes eco-trails, while a lower 'w' emphasizes electric fields.

$$P(s, d) = w * P_{eco}(s, d) + (1 - w) * P_{electric}(s, d)$$
 eq. 3

Eco-trail strengths are updated based on the energy consumption of paths taken by ants, as shown in Equation 4, 8where η represents the learning rate.

$$\Delta P_{\text{eco}}(s, d) = \eta * E(s, d)$$
 eq.

The probability of an ant moving to node d is determined by a combination of eco-trails and electric field strengths, as expressed in Equation 3, with the weight factor 'w' controlling the balance between exploration and exploitation. A higher 'w' emphasizes eco-trails, while a lower 'w' emphasizes electric fields. Eco-trail strengths are updated based on the energy consumption of paths taken by ants, as shown in Equation 4, where η represents the learning rate.

$$R_{eco}(s,d) = E_{init} - E_{cnsm}(s,d)$$
 eq. 5

Ants select paths with the highest probability to move from source node s to destination node d. The algorithm converges upon reaching a termination criterion, such as a predefined number of iterations or upon obtaining a stable/optimum route. To calculate the residual energy consumed, Equation 5 is utilized, where R_eco(s, d) represents the remaining energy at source node s after accounting for energy consumed along the path from node s to node d.

$$R_{eco}(s,d) = E_{init} - \alpha \propto D(s,d)$$
 eq. 6

Finally, Equation 6 calculates the residual energy consumed, considering the initial energy at head node s and subtracting the energy consumed along the path, thus providing the remaining energy at head node s after the energy consumed along the EcoTrial path.

Algorithm: EcoTrail
nodes:=source, destination
electric field strengths:= nodes
no. ants=X, max. iterations=N, pheromone evaporation rate=r,
convergence criteria=c





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```
While not converged and iterations < max_iterations:
  For each ant node in the network:
    Choose the destination node based on an exploration-exploitation trade-off.
    Calculate the energy consumption for the chosen path.
   end
end
  Update the eco-trails:
  For each ant_node:
    Update the strength of the eco-trail based on the energy consumption of the
chosen path.
  end
  Update the electric fields:
  For each edge (u, v) in the network:
    Calculate the electric field strength based on the charge difference between
    Update the electric field strength on edge (u, v).
Apply pheromone evaporation to eco-trails to prevent stagnation.
Check for convergence based on the specified criteria.
Choose the best path found based on eco-trails for routing.
Return the best path.
End
```

The algorithm introduces a novel method for achieving energy optimization within IoT networks, employing a graph representation with nodes (devices or sensors) and edges (connections). The approach algorithm starts by calculating the optimum route using Dijkstra's shortest path algorithm, while the weights are assigned for each edge. Further EcoTrail does the computation of energy required to be spent at each and every node (ni), during the transfer of packets (pi). The strength of the energy as well as the energy is computed. Now the presence of pheromone is verified with the chosen path, while the optimized route is selected. Once the path is chosen the pheromone count is increased with 0.1, which will be reduced in a similar manner if the path is not used for a specific duration. Figure 3, elaborates on the movement of packets which is supported by the EcoTrial algorithm. As described the process of routing starts with the preliminary assignments, followed by the computation of routes substantiated by computation of electric field strength after which the pheromone strength is verified. Passing these computations the

RESULTS AND ANALYSIS

The algorithm in question undergoes thorough testing within a simulated smart city environment facilitated by the Kooja simulator. Rigorous testing is conducted across various scenarios, including smart traffic, smart security systems, and smart waste management systems. The assessment of the algorithm's energy consumption spans multiple device configurations, starting with 10, 20, 30, and 40 sensors, respectively. Additionally, these evaluations consider the monitoring area as a pivotal parameter, analyzing energy, residual energy, and the overall performance of the algorithm. Figure 4, illustrates the amount of data generated by the devices in the G-IoT environment. Deploying 40 devices allows generation of data calculated in bits as 5000. Now considering the energy requirement for generating this huge amount of data is mandatory. Figure 5, presents the consumption of energy in watts (w) by different devices in G-IoT. Here the number of devices considered is 40. The peak representing the device in normal mode shows a maximum consumption between 150 to 400 w, in active and passive modes. On the optimization the results show significant energy savings of about 20 w to 110 w. The average consumption and savings can be





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calculated as 350 w at the normal mode and 75 w in the optimal mode. Figure 5 illustrates the residual energy consumed by the node is watts for 1700 different iterations. EcoTrial demonstrates consistent energy retention at the end of each transfer while EcoTrial performs better by the reduced consumption of energy at an average of 250 w. Though differences between energy transfer and retrieval are highly independent, variations identified show that EcoTrial outperforms the existing proposals. Figure 7 discloses the performance of the proposed EcoTrial algorithm on considering the throughput it could provide. With a varied set of inputs, the algorithm's performance is calculated in terms of packets it could transmit. It is observed that on calculating the optimal route, the throughput it provides is reliable as it provides better performance by transmitting 3.5 packets at an average, which underscores the efficiency of EcoTrial.

CONCLUSION

The study delves into the expanding realm of Green Internet of Things (G-IoT), offering a crucial response to the dynamic landscape of smart applications. It aims to harmonize technological advancements with eco-conscious practices, envisioning a future where development is not only intelligent but also inherently green and resilient. Green Internet of Things (G-IoT) applications, particularly in power management, emphasize the significance of energy efficiency. The research underscores the pivotal role of network layer resources/devices in fostering environmental friendliness. Optimization algorithms, notably Ant Colony Optimization (ACO) based schemes, substantially enhance energy consumption efficiency in IoT networks. The proposed EcoTrail algorithm, drawing inspiration from ant colony behavior and electric fields, presents a novel approach to energy optimization. It outperforms existing optimization algorithms by achieving a higher level of optimization, up to 89% while reducing the energy consumption, and providing better throughput. By amalgamating eco-trails and electric fields, the algorithm facilitates effective routing decisions, thereby promoting energy conservation and waste prevention.

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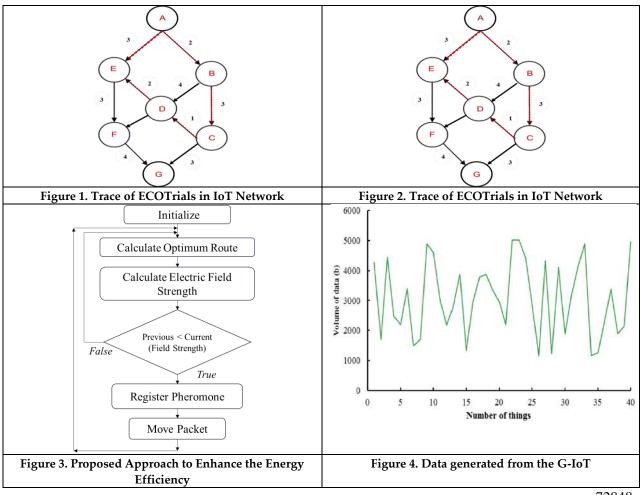
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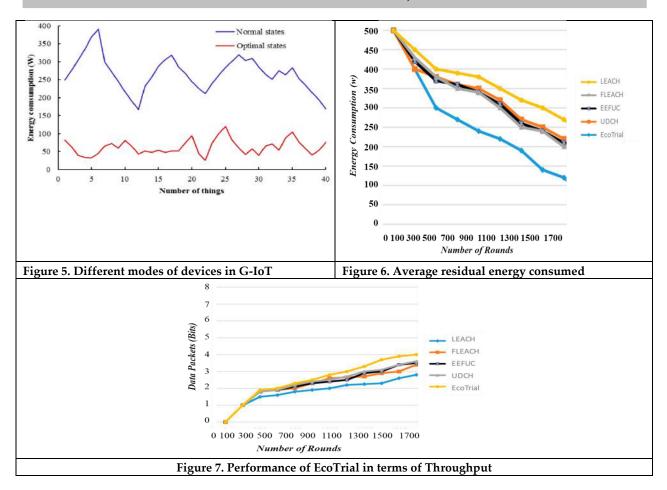
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RESEARCH ARTICLE

On Neutrosophic Λ_P - Neighbourhood and its Functions in Neutrosophic Topological Space

C. Reena^{1*} and M. Karthika²

¹Assistant Professor, Department of Mathematics, St. Mary's College (Autonomous), (Affiliated to Manonmaniam Sundaranar University, Abishekapatti, Tirunelveli) Thoothukudi, Tamil Nadu, India. ²Research Scholar, Reg.No. 21112212092004, Department of Mathematics, St. Mary's College (Autonomous), (Affiliated to Manonmaniam Sundaranar University, Abishekapatti, Tirunelveli)Thoothukudi, Tamil Nadu, India.

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*Address for Correspondence

C. Reena

Assistant Professor,

Department of Mathematics,

St. Mary's College (Autonomous),

(Affiliated to Manonmaniam Sundaranar University, Abishekapatti, Tirunelveli)

Thoothukudi, Tamil Nadu, India. Email: reenastephany@gmail.com



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ABSTRACT

The aim of this paper is to introduce the new notion of neutrosophic neighbourhood namely neutrosophic Λ P-neighbourhood and neutrosophic Λ P-quasi neighbourhood in neutrosophic topological space. Also, we introduce the neutrosophic continuous and irresolute functions along with their composition using neutrosophic Λ P-open sets.

Keywords: neutrosophic Λ_P -open, neutrosophic Λ_P -closed, neutrosophic Λ_P -neighbourhood, neutrosophic Λ_P -continuous, neutrosophic Λ_P -irresolute.

INTRODUCTION

In mathematics, Zadeh[20] introduced the concept of fuzzy set between the real standard intervals in disclipine of logic and set theory. Atanassov[3] extended the intuitionistic fuzzy set which is a combination of membership and non-membership values. Smarandache[18] created the concept of neutrosophy and neutorsophic sets at the beginning of 20th century. Later Salama[15] and Alblowi initiated the neutrosophic sets in a topology entitled as neutrosophic topological space. G. C. Ray and Sudeep[11] proposed the definitions of neutrosophic point and neighbourhood structure. They have also explored the relation of quasi coincidence between neutrosophic sets and characterized the neutrosophic topological spaces by means of quasi-neighbourhood. In 2014, Salama[16] continued the studies on neutrosophic continuous functions. Recently, the authors[12] of this paper introduced a new notion of





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neutrosophic sets namely neutrosophic Λ_P -open and neutrosophic Λ_P -closed sets. In this paper we have studied about novel concept of neutrosophic Λ_P -neighbourhood with quasi coincident. Also extended the neutrosophic continuous functions to neutrosophic Λ_P -continuous and neutrosophic Λ_P -irresolute functions with other existing continuous functions in neutrosophic topological space. Also quoted the equivalent conditions and compositions of these functions.

PRELIMINARIES

Definition 2.1:[15]Let U be a non-empty fixed set. A **Neutrosophic set**K is an object having the form $K = \{\langle u, \mu_K(u), \sigma_K(u), \gamma_K(u) \rangle: u \in U\}$ where $\mu_K(u), \sigma_K(u)$ and $\gamma_K(u)$ represents the degree of membership, the degree of indeterminacy and the degree of non-membership respectively of each element $u \in U$ to the set U. A neutrosophic set $K = \{\langle u, \mu_K(u), \sigma_K(u), \gamma_K(u) \rangle: u \in U\}$ can be identified to an ordered triple $\langle \mu_K(u), \sigma_K(u), \gamma_K(u) \rangle$ in $1^{-0.1^{+}}$ on U.

Definition 2.2:[15]Let U be a non-empty set and $K = \{\langle u, \mu_K(u), \sigma_K(u), \gamma_K(u) \rangle : u \in U\}$ and $M = \{\langle u, \mu_M(u), \sigma_M(u), \gamma_M(u) \rangle : u \in U\}$ are neutrosophic sets, then

- i. $K \subseteq M \Leftrightarrow \mu_K(u) \le \mu_M(u), \sigma_K(u) \le \sigma_M(u) \text{ and } \gamma_K(u) \ge \gamma_M(u) \forall u \in U$
- ii. $K \cup M = \{(u, \max(\mu_K(u), \mu_M(u)), \max(\sigma_K(u), \sigma_M(u)), \min(\gamma_K(u), \gamma_M(u)) : u \in U)\}$
- iii. $K \cap M = \{(u, \min(\mu_K(u), \mu_M(u)), \min(\sigma_K(u), \sigma_M(u)), \max(\gamma_K(u), \gamma_M(u)) : u \in U\}\}$
- iv. $K^C = \{\langle u, (\gamma_K(u), 1 \sigma_K(u), \mu_K(u)) \rangle : u \in U \}$
- v. $0_{N_{tr}} = \{\langle u, 0, 0, 1 \rangle : u \in U\} \text{ and } 1_{N_{tr}} = \{\langle u, 1, 1, 0 \rangle : u \in U\}$

Definition 2.3:[15]A **Neutrosophic topology** on a non-empty set U is a family $\tau_{N_{tr}}$ of neutrosophic sets in U satisfying the following axioms:

- i. $0_{N_{tr}}, 1_{N_{tr}} \in \tau_{N_{tr}}$.
- ii. $K_1 \cap K_2 \in \tau_{N_{tr}}$ for any $K_1, K_2 \in \tau_{N_{tr}}$.
- iii. $\bigcup K_i \in \tau_{N_{tr}} \text{ for every } \{K_i : i \in I\} \subseteq \tau_{N_{tr}}.$

In this case the ordered pair $(U, \tau_{N_{tr}})$ is called a neutrosophic topological space. The members of $\tau_{N_{tr}}$ are neutrosophic open set and its complements are neutrosophic closed.

Definition 2.4:[6]A neutrosophic set $K = \{\langle u, \mu_K(u), \sigma_K(u), \gamma_K(u) \rangle : u \in U \}$ is called a **neutrosophic point** if for any element $v \in U$, $\mu_K(v) = a$, $\sigma_K(v) = b$, $\gamma_K(v) = c$ and for u = v and $\mu_K(v) = 0$, $\sigma_K(v) = 0$, $\gamma_K(v) = 1$ for $u \neq v$, where a, b, c are real standard or non standard sunsets of . A neutrosophic point is denoted by $u_{a,b,c}$. For the neutrosophic point $u_{a,b,c}$, u will be called its support.

Definition 2.5[10]: Let $u_{a,b,c}$ be a neutrosophic point in a neutrosophic topological space $(U, \tau_{N_{tr}})$. Then a neutrosophic set N in U is said to be **neutrosophicneighbourhood**(N_{tr} nbhd) of $u_{a,b,c}$ if there exists a N_{tr} open set M such that $u_{a,b,c} \in M \subseteq N$.

Definition 2.6[1]: A neutrosophic point $u_{a,b,c}$ is said to be **neutrosophic quasi - coincident** with a neutrosophic set K, denoted by $u_{a,b,c}qK$ if $u_{a,b,c} \notin K^c$. If $u_{a,b,c}$ is not neutrosophic quasi – coincident with K, we denote it by $u_{a,b,c}\hat{q}K$.

Definition 2.7[1]: A neutrosophic set M is said to be neutrosophic quasi - coincident with a neutrosophic set K, denoted by MqK if $M \nsubseteq K^c$. If M is not neutrosophic quasi - coincident with K, we denote it by $M\hat{q}K$.

Definition 2.8[1]: A neutrosophic set N in U is said to be **neutrosophic quasi- neighbourhood**(N_{tr} Qnbhd) of $u_{a,b,c}$ if there exists a N_{tr} open set M such that $u_{a,b,c}$ $qM \subseteq N$.

Definition 2.9[8]: Let $(U, \tau_{N_{tr}})$ be a neutrosophic topological space and S be a non-empty subset of U. Then, a neutrosophic relative topology on S is defined by





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 $\tau_{N_{tr}}^S = \{K \cap \mathbb{1}_{N_{tr}}^S : K \in \tau_{N_{tr}}\}$ where

$$1_{N_{tr}}^{S} = \begin{cases} <1,1,0>, & if s \in S\\ <0,0,1>, & otherwise \end{cases}$$
space of $(U, \tau_{N_{tr}})$.

Thus, $(S, \tau_{N_{tr}}^{S})$ is called a neutrosophic subspace of $(U, \tau_{N_{tr}})$.

Definition 2.10:[5] Let U and V be two non-empty sets and $f_{N_{tr}}: U \to V$ be a function. If $M = \{\langle v, \mu_M(v), \sigma_M(v), \gamma_M(v) \rangle : v \in V\}$ is a neutrosophic set in V, then the **pre-image** of M under

under $f_{N_{tr}}$ denoted by $f_{N_{tr}}^{-1}(M)$, is the neutrosophic set in U defined by

$$f_{N_{tr}}^{-1}(M) = \{ \langle u, f_{N_{tr}}^{-1}(\mu_M)(u), f_{N_{tr}}^{-1}(\sigma_M)(u), f_{N_{tr}}^{-1}(\gamma_M)(u) \rangle : u \in U \}$$

If $K = \{\langle u, \mu_K(u), \sigma_K(u), \gamma_K(u) \rangle : u \in U\}$ is a neutrosophic set in U, then the **image** of K under $f_{N_{tr}}$, denoted by $f_{N_{tr}}(K)$, is the neutrosophic set in V defined by

$$f_{N_{tr}}(K) = \{ \langle v, f_{N_{tr}}(\mu_K)(v), f_{N_{tr}}(\sigma_K)(v), \left(1 - f_{N_{tr}}(1 - \gamma_K)\right)(v) \rangle : v \in V \} \text{where}$$

$$f_{N_{tr}}(\mu_K)(v) = \begin{cases} \sup_{u \in f_{N_{tr}}^{-1}(v)} \mu_K(u), & \text{if } f_{N_{tr}}^{-1}(v) \neq \emptyset \\ 0, & \text{otherwise} \end{cases}$$

$$f_{N_{tr}}(\sigma_K)(v) = \begin{cases} \sup_{u \in f_{N_{tr}}^{-1}(v)} \sigma_K(u), & \text{if } f_{N_{tr}}^{-1}(v) \neq \emptyset \\ 0, & \text{otherwise} \end{cases}$$

$$\left(1 - f_{N_{tr}}(1 - \gamma_K)\right)(v) = \begin{cases} \inf_{u \in f_{N_{tr}}^{-1}(v)} \gamma_K(u), & \text{if } f_{N_{tr}}^{-1}(v) \neq \emptyset \\ 1, & \text{otherwise} \end{cases}$$

Definition 2.11: Let $(U, \tau_{N_{tr}})$ and $(V, \rho_{N_{tr}})$ be neutrosophic topological spaces. Then the function $f_{N_{tr}}: (U, \tau_{N_{tr}}) \to (V, \rho_{N_{tr}})$ is said to be **neutrosophic continuous**[15] (respectively, neutrosophic semi-continuous[4], neutrosophic α-continuous[2], neutrosophic regular-continuous[19], neutrosophic δ-continuous, neutrosophic b-continuous, neutrosophic Λ_p^* -continuous) if $f_{N_{tr}}^{-1}(M)$ is N_{tr} open(respectively N_{tr} semi-open, N_{tr} α-open, N_{tr} α-open, N_{tr} δ-open, N_{tr} δ-open, N_{tr} δ-open, N_{tr} Λ_p^* -open) in $(U, \tau_{N_{tr}})$ for every N_{tr} open set M in $(V, \rho_{N_{tr}})$.

Definition 2.12:[12]A neutrosophic set K of a neutrosophic topological space $(U, \tau_{N_{tr}})$ is said to be **neutrosophic** Λ_{P} -**open** if there exist a N_{tr} pre-open set $E \neq 0_{N_{tr}}, 1_{N_{tr}}$ such that $K \subseteq N_{tr}$ $cl(K \cap E)$. The complement of neutrosophic Λ_{P} open set is neutrosophic Λ_{P} -closed. The class of neutrosophic Λ_{P} -open sets is denoted by $N_{tr}\Lambda_{P}O(U, \tau_{N_{tr}})$.

Theorem 2.13:[12] The union of an arbitrary collection of $N_{tr} \Lambda_{P}$ -open sets is also $N_{tr} \Lambda_{P}$ -open.

Theorem 2.14:[12]In any neutrosophic topological space $(U, \tau_{N_{tr}})$,

- i. Every N_{tr} -open set is $N_{tr} \Lambda_{P}$ -open.
- ii. Every N_{tr} regular-open set is $N_{tr} \Lambda_{P}$ -open.
- iii. Every $N_{tr} \delta$ -open set is $N_{tr} \Lambda_{P}$ -open.
- iv. Every N_{tr} semi-open set is $N_{tr} \Lambda_P$ -open.
- v. Every $N_{tr} \alpha$ -open set is $N_{tr} \Lambda_{P}$ -open.
- vi. Every $N_{tr} \Lambda_{P}$ -open set is $N_{tr} \Lambda_{P}^{*}$ -open.

Remark 2.15:[12] The above theorem is also true for $N_{tr} \Lambda_P$ -closed sets.

Theorem 2.16:[12]A neutrosophic set K in a neutrosophic topological space $(U, \tau_{N_{tr}})$ is $N_{tr} \Lambda_{P}$ -open if and only if for every neutrosophic point $u_{a,b,c} \in K$, there exists a $N_{tr} \Lambda_{P}$ -open set $M_{u_{a,b,c}}$ such that $u_{a,b,c} \in M_{u_{a,b,c}} \subseteq K$.

Definition 2.17:[12]Let $(U, \tau_{N_{tr}})$ be a neutrosophic topological space and K be a neutrosophic set in U.

i. The **neutrosophic** Λ_P -interior of K is the union of all $N_{tr}\Lambda_P$ -open sets contained in K. It is denoted by $N_{tr}\Lambda_P$ int(K).





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ii. The **neutrosophic** $\Lambda_{\mathbf{P}}$ -**closure** of K is the intersection of all N_{tr} -closed sets containing K. It is denoted by $N_{tr} \Lambda_{\mathbf{P}} cl(K)$.

Theorem 3.10: A neutrosophic set K in a neutrosophic topological space $(U, \tau_{N_{tr}})$ is $N_{tr} \Lambda_{P}$ -open if and only if for every neutrosophic point $u_{a,b,c} \in K$, K is a $N_{tr} \Lambda_{P}$ -nbhd of $u_{a,b,c}$.

Proof: Let K be $N_{tr}\Lambda_P$ -open in U. Then for each $u_{a,b,c} \in K$, $K \subseteq K$. Therefore K is a $N_{tr}\Lambda_P$ -nbhd of $u_{a,b,c}$. Conversely, assume that for every $u_{a,b,c} \in K$, K is a $N_{tr}\Lambda_P$ -nbhd of $u_{a,b,c}$. Then, there exists a $N_{tr}\Lambda_P$ -open set K in U such that $u_{a,b,c} \in M \subseteq K$. Therefore, by theorem 2.16, K is $N_{tr}\Lambda_P$ -open.

Theorem 3.11: Every $N_{tr}\Lambda_P$ -open set K in a neutrosophic topological space $(U, \tau_{N_{tr}})$ is a $N_{tr}\Lambda_P$ -Qnbhd of every neutrosophic point quasi-coincident with K.

Proof: The proof is obvious since for every neutrosophic point $u_{a,b,c}qK$, we have $u_{a,b,c}qK \subseteq K$.

Theorem 3.12: Let K be a $N_{tr}\Lambda_P$ -closed set in a neutrosophic topological space $(U, \tau_{N_{tr}})$ and $u_{a,b,c}qK^C$. Then there exists a $N_{tr}\Lambda_P$ -QnbhdM of $u_{a,b,c}$ such that $K\hat{q}M$.

Proof: Since K is $N_{tr} \Lambda_P$ -closed in U, K^C is $N_{tr} \Lambda_P$ -open in U such that $u_{a,b,c} q K^C$. Then, by theorem 3.11, K^C is a $N_{tr} \Lambda_P$ -Qnbhd of $u_{a,b,c}$. Hence there exists $aN_{tr} \Lambda_P$ -open set M in Usuch that $u_{a,b,c} q K \subseteq K^C$ Again, by theorem 3.11, M is a $N_{tr} \Lambda_P$ -Qnbhd of $u_{a,b,c}$ Also, since $M \subseteq K^C$, $K \widehat{q} M$.

NEUTROSOPHIC Λ_P-NEIGHBOURHOOD

Definition 3.1: Let $u_{a,b,c}$ be a neutrosophic point in a neutrosophic topological space $(U, \tau_{N_{tr}})$. Then a neutrosophic set N in U is said to be

- i. **neutrosophic** Λ_{P} -**neighbourhood**($N_{tr}\Lambda_{P}$ -nbhd) of $u_{a,b,c}$ if there exists a $N_{tr}\Lambda_{P}$ -open set M such that $u_{a,b,c} \in M \subseteq N$.
- ii. **neutrosophic** Λ_{P} **-quasi neighbourhood**($N_{tr}\Lambda_{P}$ -Qnbhd) of $u_{a,b,c}$ if there exists a $N_{tr}\Lambda_{P}$ -open set M such that $u_{a,b,c}qM \subseteq N$.

Example 3.2: Let $U = \{a, b\}$ and $\tau_{N_{tr}} = \{0_{N_{tr}}, 1_{N_{tr}}, K\}$ where $K = \{< a, 0.6, 0.4, 0.2 > < b, 0.1, 0.6, 0.1 > \}$. Now, let us consider a neutrosophic point $a_{0.1,0.2,0.4}$ in U. Then there exists a $N_{tr} \Lambda_{P}$ -open set $M = \{< a, 0.7, 0.6, 0.1 > < b, 0.4, 0.8, 0.1 > \}$ such that $a_{0.1,0.2,0.4} \in M \subseteq N$ where $N = \{< a, 0.8, 0.7, 0.1 > < b, 0.5, 0.8, 0.1 > \}$. Hence N is a $N_{tr} \Lambda_{P}$ -nbhd of $a_{0.1,0.2,0.4}$.

Example 3.3: Let $U = \{a, b\}$ and $\tau_{N_{tr}} = \{0_{N_{tr}}, 1_{N_{tr}}, K\}$ where $K = \{< a, 0.7, 0.6, 0.1 >< b, 0.4, 0.8, 0.1 >\}$. Now, let us consider a neutrosophic point $a_{0.1,0.8,0.7}$ in U. Then there exists a $N_{tr} \Lambda_{P}$ -open set $M = \{< a, 0.8, 0.7, 0.1 >< b, 0.7, 0.8, 0.1 >\}$ such that $a_{0.1,0.8,0.7} qM \subseteq N$ where $N = \{< a, 0.8, 0.8, 0.8, 0.1 >< b, 0.7, 0.8, 0.1 >\}$. Hence N is a $N_{tr} \Lambda_{P}$ -Qnbhd of $a_{0.1,0.8,0.7}$.

Theorem 3.4: Every N_{tr} nbhd of a neutrosophic point $u_{a,b,c}$ in a neutrosophic topological space $(U, \tau_{N_{tr}})$ is a $N_{tr} \Lambda_{P}$ nbhd of $u_{a,b,c}$.

Proof: Let N be a N_{tr} nbhd of a neutrosophic point $u_{a,b,c}$ in U. Then, there exists a N_{tr} open set M in U such that $u_{a,b,c} \in M \subseteq N$. Now, by theorem 2.14, M is $N_{tr}\Lambda_P$ -open in U. Hence there exists a $N_{tr}\Lambda_P$ -open set M in U such that $u_{a,b,c} \in M \subseteq N$. Therefore N is a $N_{tr}\Lambda_P$ -nbhd of $u_{a,b,c}$.

Remark 3.5: The above theorem's converse need not hold.

Example 3.6:Let $U = \{a, b\}$ and $\tau_{N_{tr}} = \{0_{N_{tr}}, 1_{N_{tr}}, K\}$ where $K = \{< a, 0.5, 0.5, 0.1 >< b, 0.1, 0.8, 0.1 >\}$. Now, let us consider a neutrosophic point $a_{0.6, 0.1, 0.4}$ in U. Then there exists a $N_{tr} \Lambda_P$ -open set





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 $M = \{ < a, 0.7, 0.7, 0.1 > < b, 0.2, 0.8, 0.1 > \}$ such that $a_{0.6,0.1,0.4} \in M \subseteq N$ where $N = \{ < a, 0.7, 0.8, 0.1 > < b, 0.3, 0.8, 0.1 > \}$. This implies N is a $N_{tr} \Lambda_P$ -nbhd of $a_{0.6,0.1,0.4}$. However, N is not a N_{tr} nbhd of $a_{0.6,0.1,0.4}$.

Theorem 3.7: Every N_{tr} Qnbhdof a neutrosophic point $u_{a,b,c}$ in a neutrosophic topological space $(U, \tau_{N_{tr}})$ is $aN_{tr} \Lambda_{P}$ -Qnbhd of $u_{a,b,c}$.

Proof: Let N be a N_{tr} Qnbhdof a neutrosophic point $u_{a,b,c}$ in U. Then, there exists a N_{tr} open set M in U such that $u_{a,b,c}$ $qM \subseteq N$. Now, by theorem 2.14, M is $N_{tr} \Lambda_P$ -open in U. Hence there exists a $N_{tr} \Lambda_P$ -open set M in U such that $u_{a,b,c}$ $qM \subseteq N$. Thus N is a $N_{tr} \Lambda_P$ -Qnbhd of $u_{a,b,c}$.

Remark 3.8: The above theorem's converse need not hold.

Example 3.9: Let $U = \{a, b\}$ and $\tau_{N_{tr}} = \{0_{N_{tr}}, 1_{N_{tr}}, K\}$ where $K = \{< a, 0.6, 0.8, 0.1 > < b, 0.4, 0.6, 0.3 > \}$. Now, let us consider a neutrosophic point $a_{0.1,0.1,0.5}$ in U. Then there exists a $N_{tr}\Lambda_{P}$ -open set $M = \{< a, 0.7, 0.8, 0.1 > < b, 0.6, 0.6, 0.6, 0.1 > \}$ such that $a_{0.1,0.1,0.5}qM \subseteq N$ where $N = \{< a, 0.8, 0.8, 0.8, 0.1 > < a, 0.8, 0.7, 0.1 > \}$. This implies N is a $N_{tr}\Lambda_{P}$ -Qnbhd of $a_{0.1,0.1,0.5}$. However, N is not a $N_{tr}\Lambda_{P}$ -Qnbhd of $a_{0.1,0.1,0.5}$.

NEUTROSOPHIC Λ_P -CONTINUOUS FUNCTIONS

Definition 4.1: Let $(U, \tau_{N_{tr}})$ and $(V, \rho_{N_{tr}})$ be neutrosophic topological spaces. Then the function $f_{N_{tr}}: (U, \tau_{N_{tr}}) \to (V, \rho_{N_{tr}})$ is said to be **neutrosophic** Λ_P -**continuous** if $f_{N_{tr}}^{-1}(M)$ is $N_{tr}\Lambda_P$ -open in $(U, \tau_{N_{tr}})$ for every N_{tr} -open set M in $(V, \rho_{N_{tr}})$.

Example 4.2: Let $U = \{a, b\}, V = \{x, y\}, \tau_{N_{tr}} = \{0_{N_{tr}}, 1_{N_{tr}}, K, M\}$ and $\rho_{N_{tr}} = \{0_{N_{tr}}, 1_{N_{tr}}, N\}$ where $K = \{< a, 0.7, 0.8, 0.4 > < b, 1, 1, 0 >, M = \{< a, 0.5, 0.6, 0.5 > < b, 0.6, 0.8, 0.1 > \}$ and $N = \{< x, 0.8, 0.9, 0.1 > < y, 1, 1, 0 > \}$. Consider the collections $A = \{A, K \subset A, M^c \subset A\}$ and $B = \{B : K \subset B; B \not\subset M^c; M^c \not\subset B\}$ of neutrosophic sets in U. Then $N_{tr}\Lambda_P O(U, \tau_{N_{tr}}) = \{0_{N_{tr}}, K, M, A, B, 1_{N_{tr}}\}$. Define $f_{N_{tr}} : (U, \tau_{N_{tr}}) \to (V, \rho_{N_{tr}})$ as $f_{N_{tr}}(a) = y$ and $f_{N_{tr}}(b) = x$. Then, $f_{N_{tr}}^{-1}(N) = \{< a, 1, 1, 0 > < b, 0.8, 0.9, 0.1 > \} \in A$ which implies $f_{N_{tr}}^{-1}(N)$ is $N_{tr}\Lambda_P$ -open in U. Hence $f_{N_{tr}}$ is $N_{tr}\Lambda_P$ -continuous

Theorem 4.3: Every N_{tr} -continuous function is $N_{tr}\Lambda_P$ -continuous.

Proof: Let $f_{N_{tr}}: (U, \tau_{N_{tr}}) \to (V, \rho_{N_{tr}})$ be a N_{tr} -continuous function. Let M be a N_{tr} -open set in V. Since $f_{N_{tr}}$ is N_{tr} -continuous, $f_{N_{tr}}^{-1}(M)$ is N_{tr} -open in U. Hence $f_{N_{tr}}$ is $N_{tr}\Lambda_{P}$ -continuous.

Remark 4.4: The above theorem's converse need not hold.

Example 4.5: Let $U = \{a, b\}, V = \{x, y\}, \tau_{N_{tr}} = \{0_{N_{tr}}, K, M, 1_{N_{tr}}\}$ and $\rho_{N_{tr}} = \{0_{N_{tr}}, N, 1_{N_{tr}}\}$ where $K = \{< a, 0.3, 0.4, 0.9 > < b, 0.4, 0.5, 0.8 > \}, M = \{< a, 0.2, 0.3, 1 > < b, 0.3, 0.4, 0.9 > \}$ and $N = \{< x, 0.6, 0.5, 0.3 > < y, 0.7, 0.6, 0.2 > \}$. Consider the collections $A = \{A : M \subset A \subset K^c\}$ and $B = \{B : B \subset M^c; B \not\subset K; K \not\subset B\}$ of neutrosophic sets in U. Then $N_{tr}\Lambda_P O(U, \tau_{N_{tr}}) = \{0_{N_{tr}}, K, M, A, B, 1_{N_{tr}}\}$. Define $f_{N_{tr}}: (U, \tau_{N_{tr}}) \to (V, \rho_{N_{tr}})$ as $f_{N_{tr}}(a) = y$ and $f_{N_{tr}}(b) = x$. Then, $f_{N_{tr}}^{-1}(N) = \{< a, 0.7, 0.6, 0.2 > < b, 0.6, 0.5, 0.3 > \} ∈ Q$ which implies $f_{N_{tr}}^{-1}(N)$ is $N_{tr}\Lambda_P$ -open but not N_{tr} open in U. Hence $f_{N_{tr}}$ is $N_{tr}\Lambda_P$ -continuous but not N_{tr} -continuous.

Theorem 4.6: Let $f_{N_{tr}}:(U,\tau_{N_{tr}}) \to (V,\rho_{N_{tr}})$ be a function between two neutrosophic topological spaces.

- i. If $f_{N_{tr}}$ is N_{tr} regular-continuous, then $f_{N_{tr}}$ is $N_{tr}\Lambda_P$ -continuous.
- ii. If $f_{N_{tr}}$ is $N_{tr}\delta$ -continuous, then $f_{N_{tr}}$ is $N_{tr}\Lambda_P$ -continuous.
- iii. If $f_{N_{tr}}$ is N_{tr} semi-continuous, then $f_{N_{tr}}$ is $N_{tr}\Lambda_P$ -continuous.
- iv. If $f_{N_{tr}}$ is $N_{tr}\alpha$ -continuous, then $f_{N_{tr}}$ is $N_{tr}\Lambda_P$ -continuous.
- v. If $f_{N_{tr}}$ is $N_{tr}\Lambda_P$ -continuous, then $f_{N_{tr}}$ is $N_{tr}\Lambda_P^*$ -continuous.

Proof: Proof is obvious.

Remark 4.7: The above statements converse need not hold.





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Example 4.8: Let $U = \{a, b\}, V = \{x, y\}, \tau_{N_{tr}} = \{0_{N_{tr}}, K, 1_{N_{tr}}\} \text{and} \rho_{N_{tr}} = \{0_{N_{tr}}, M, 1_{N_{tr}}\} \text{ where } K = \{< a, 0.7, 0.6, 0.8 > < b, 0.6, 0.8, 0.9 >, <math>M = \{< a, 0.6, 0.4, 0.9 > < b, 0.5, 0.7, 0.1 > \}$. Consider the collections $\mathcal{A} = \{A : K \subset A; A^C \subset K\}, \mathcal{B} = \{B : K \subset B; B \not\subset K^C\}$ of neutrosophic sets in U. Then, $N_{tr}RO(U, \tau_{N_{tr}}) = \{0_{N_{tr}}, 1_{N_{tr}}\}$, and $N_{tr}\Lambda_PO(U, \tau_{N_{tr}}) = \{0_{N_{tr}}, K, \mathcal{A}, \mathcal{B}, 1_{N_{tr}}\}$. Define $f_{N_{tr}}: (U, \tau_{N_{tr}}) \to (V, \rho_{N_{tr}})$ as $f_{N_{tr}}(a) = y$ and $f_{N_{tr}}(b) = x$. Then, $f_{N_{tr}}^{-1}(M) = \{< a, 0.5, 0.7, 0.1 > < b, 0.6, 0.4, 0.9 > \} ∈ \mathcal{B}$ which implies $f_{N_{tr}}^{-1}(M)$ is $N_{tr}\Lambda_{P}$ -open but not N_{tr} regular-open in U. Hence $f_{N_{tr}}$ is $N_{tr}\Lambda_{P}$ -continuous but not N_{tr} regular-continuous.

Example 4.9: Let $U = \{a, b\}, V = \{x, y\}, \tau_{N_{tr}} = \{0_{N_{tr}}, K, 1_{N_{tr}}\} \text{and} \rho_{N_{tr}} = \{0_{N_{tr}}, M, 1_{N_{tr}}\}$ where $K = \{< a, 0.4, 0.8, 0.3 > < b, 0.5, 0.7, 0.2 >, <math>M = \{< a, 0.5, 0.8, 0.3 > < b, 0.2, 0.7, 0.4 >\}$. Consider the collections $\mathcal{A} = \{A : K^{C} \subset A \subset A\}, \mathcal{B} = \{B : K \subset B \subset 1_{N_{tr}}\} \text{of}$ neutrosophic sets in U. Then, $N_{tr} \delta O(U, \tau_{N_{tr}}) = \{0_{N_{tr}}, K, 1_{N_{tr}}\} \text{and} N_{tr} \Lambda_{P} O(U, \tau_{N_{tr}}) = \{0_{N_{tr}}, K, \mathcal{A}, \mathcal{B}, 1_{N_{tr}}\}.$ Define $f_{N_{tr}} : (U, \tau_{N_{tr}}) \to (V, \rho_{N_{tr}})$ as $f_{N_{tr}}(a) = y$ and $f_{N_{tr}}(b) = x$. Then, $f_{N_{tr}}^{-1}(M) = \{< a, 0.2, 0.7, 0.4 > < b, 0.5, 0.8, 0.3 > \} ∈ \mathcal{A}$ which implies $f_{N_{tr}}^{-1}(M)$ is $N_{tr} \Lambda_{P}$ -open but not $N_{tr} \delta$ -open in U. Hence $f_{N_{tr}}$ is $N_{tr} \Lambda_{P}$ -continuous but not $N_{tr} \delta$ -continuous.

Example 4.10: Let $U = \{a,b\}, V = \{x,y\}, \tau_{N_{tr}} = \{0_{N_{tr}}, K, 1_{N_{tr}}\}$ and $\rho_{N_{tr}} = \{0_{N_{tr}}, M, 1_{N_{tr}}\}$ where $K = \{< \alpha, 0.4, 0.5, 0.6 > < b, 0.6, 0.3, 0.8 >, <math>M = \{< x, 0.3, 0.4, 0.7 > < y, 0.5, 0.2, 0.9 > \}$. Consider the collections $\mathcal{A} = \{A : 0_{N_{tr}} \subset A \subset K\}, \mathcal{B} = \{B : K^{C} \subset B \subset 1_{N_{tr}}\}$ and $\mathcal{C} = \{C : K \subset C \not\subset K^{C}\}$ of neutrosophic sets in U. Then, $N_{tr}\alpha O(U, \tau_{N_{tr}}) = \{0_{N_{tr}}, 1_{N_{tr}}, K, M\}, N_{tr}SO(U, \tau_{N_{tr}}) = \{0_{N_{tr}}, K, M, K^{C}, M^{C}, 1_{N_{tr}}\}$ and $N_{tr}\Lambda_{P}O(U, \tau_{N_{tr}}) = \{0_{N_{tr}}, M, M^{C}, \mathcal{A}, \mathcal{B}, \mathcal{C}, 1_{N_{tr}}\}$. Define $f_{N_{tr}}: (U, \tau_{N_{tr}}) \to (V, \rho_{N_{tr}})$ as $f_{N_{tr}}(a) = y$ and $f_{N_{tr}}(b) = x$. Then, $f_{N_{tr}}^{-1}(M) = \{< \alpha, 0.5, 0.2, 0.9 > < b, 0.3, 0.4, 0.7 > \} ∈ \mathcal{A}$ which implies $f_{N_{tr}}^{-1}(M)$ is $N_{tr}\Lambda_{P}$ -open. However, it is neither N_{tr} semi-open nor $N_{tr}\alpha$ -open in U. Hence $f_{N_{tr}}$ is $N_{tr}\Lambda_{P}$ -continuous but not N_{tr} semi-continuous and $N_{tr}\alpha$ -continuous.

Example 4.11: Let $U = \{a,b\}, V = \{x,y\}, \tau_{N_{tr}} = \{0_{N_{tr}}, 1_{N_{tr}}, K\}$ and $\rho_{N_{tr}} = \{0_{N_{tr}}, 1_{N_{tr}}, M\}$ where $K = \{< a, 0.7, 0.6, 0.8 > < b, 0.6, 0.8, 0.9 > \}$ and $M = \{< x, 0.5, 0.7, 0.2 > < y, 0.6, 0.9, 0.1 > \}$. Consider the collections $\mathcal{A} = \{A : K^{\mathcal{C}} \subset A \subset K\}, \mathcal{B} = \{B : K \subset B \subset 1_{N_{tr}}\}, \mathcal{C} = \{C : K^{\mathcal{C}} \not\subset \mathcal{C} \not\subset K^{\mathcal{C}}; \mathcal{C} \subset K\}, \mathcal{D} = \{D : K^{\mathcal{C}} \not\subset D; D \not\subset K^{\mathcal{C}}; \mathcal{D} \not\subset K\}, \mathcal{E} = \{E : K^{\mathcal{C}} \subset E \not\subset K\} \text{ and } \mathcal{F} = \{F : 0_{N_{tr}} \subset F \subset K^{\mathcal{C}}\} \text{ of neutrosophic sets in } U.$ Then, $N_{tr}\Lambda_{P}^{*}0(U, \tau_{N_{tr}}) = \{0_{N_{tr}}, K, \mathcal{A}, \mathcal{B}, \mathcal{C}, \mathcal{D}, \mathcal{E}, \mathcal{F}, 1_{N_{tr}}\},$ and $N_{tr}\Lambda_{P}0(U, \tau_{N_{tr}}) = \{0_{N_{tr}}, K, \mathcal{B}, \mathcal{F}, 1_{N_{tr}}\}.$ Define $f_{N_{tr}}: (U, \tau_{N_{tr}}) \to (V, \rho_{N_{tr}}) \text{ as } f_{N_{tr}}(a) = x \text{ and } f_{N_{tr}}(b) = y.$ Then, $f_{N_{tr}}^{-1}(M) = \{< a, 0.5, 0.7, 0.2 > < b, 0.6, 0.9, 0.1 > \} \in \mathcal{D}$ which implies $f_{N_{tr}}^{-1}(M)$ is $N_{tr}\Lambda_{P}^{*}$ -open, but not $N_{tr}\Lambda_{P}$ -open. Hence $f_{N_{tr}}$ is $N_{tr}\Lambda_{P}^{*}$ -continuous, but not $N_{tr}\Lambda_{P}$ -continuous.

Theorem 4.12: Let $f_{N_{tr}}:(U,\tau_{N_{tr}}) \to (V,\rho_{N_{tr}})$ be a function between two neutrosophic topological spaces. Then the following statements are equivalent:

- i. $f_{N_{tr}}$ is $N_{tr}\Lambda_P$ -continuous.
- ii. The inverse image of every N_{tr} -closed set in $(V, \rho_{N_{tr}})$ is $N_{tr} \Lambda_P$ -closed in $(U, \tau_{N_{tr}})$.
- iii. $f_{N_{tr}}(N_{tr}\Lambda_P cl(K)) \subseteq N_{tr} cl(f_{N_{tr}}(K))$ for every neutrosophic set K in U.
- iv. $N_{tr}\Lambda_P cl\left(f_{N_{tr}}^{-1}(M)\right)\subseteq f_{N_{tr}}^{-1}\left(N_{tr}cl(M)\right)$ for every neutrosophic set M in V.

Proof:

(i) \Rightarrow (ii) Let $f_{N_{tr}}$ be a $N_{tr}\Lambda_P$ -continuous function and M be a N_{tr} closed set in V. Then M^c is N_{tr} open in V. Since $f_{N_{tr}}$ is $N_{tr}\Lambda_P$ -continuous, $f_{N_{tr}}^{-1}(M^C)$ is $N_{tr}\Lambda_P$ -open in U which implies $(f_{N_{tr}}^{-1}(M))^C$ is $N_{tr}\Lambda_P$ -open in U. Hence $f_{N_{tr}}^{-1}(M)$ is $N_{tr}\Lambda_P$ -closed in U.

(ii) \Rightarrow (i) Let M be N_{tr} open in V. Then M^C is N_{tr} closed in V. By assumption, $f_{N_{tr}}^{-1}(M^C)$ is $N_{tr}\Lambda_P$ -closed in U. That is, $(f_{N_{tr}}^{-1}(M))^C$ is $N_{tr}\Lambda_P$ -closed in U. Hence $f_{N_{tr}}^{-1}(M)$ is $N_{tr}\Lambda_P$ -open in U. Therefore, $f_{N_{tr}}$ is $N_{tr}\Lambda_P$ -continuous.

(ii) \Rightarrow (iii) Let K be a neutrosophic set in U. Now, $K \subseteq f_{N_{tr}}^{-1}\left(f_{N_{tr}}(K)\right)$ implies $K \subseteq f_{N_{tr}}^{-1}\left(N_{tr}cl\left(f_{N_{tr}}(K)\right)\right)$. Since $N_{tr}cl\left(f_{N_{tr}}(K)\right)$ is N_{tr} closed in V, by assumption $f_{N_{tr}}^{-1}\left(N_{tr}cl\left(f_{N_{tr}}(K)\right)\right)$ is a $N_{tr}\Lambda_{P}$ -closed set containing K. Also,





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 $N_{tr}\Lambda_P cl(K)$ is the smallest $N_{tr}\Lambda_P$ -closed set containing K. Hence, $N_{tr}\Lambda_P cl(K) \subseteq f_{N_{tr}}^{-1}(N_{tr}cl(f_{N_{tr}}(K)))$. Therefore, $f_{N_{tr}}(N_{tr}\Lambda_P cl(K)) \subseteq N_{tr}cl(f_{N_{tr}}(K))$.

(iii) \Rightarrow (ii) Let *M* be a N_{tr} closed set in *V*. Then, by assumption

 $f_{N_{tr}}\left(N_{tr}\Lambda_{P}cl\left(f_{N_{tr}}^{-1}(M)\right)\right)\subseteq N_{tr}cl\left(f_{N_{tr}}\left(f_{N_{tr}}^{-1}(M)\right)\right)\subseteq N_{tr}cl(M)=M \text{ which implies } N_{tr}\Lambda_{P}cl\left(f_{N_{tr}}^{-1}(M)\right)\subseteq f_{N_{tr}}^{-1}(M). \text{ Also,}$ $f_{N_{tr}}^{-1}(M)\subseteq N_{tr}\Lambda_{P}cl\left(f_{N_{tr}}^{-1}(M)\right). \text{ Hence } f_{N_{tr}}^{-1}(M) \text{ is } N_{tr}\Lambda_{P}\text{-closed in } U.$

(iii) \Rightarrow (iv) Let M be a neutrosophic set in V and let $K = f_{N_{tr}}^{-1}(M)$. By assumption, $f_{N_{tr}}(N_{tr}\Lambda_P cl(K)) \subseteq N_{tr} cl(f_{N_{tr}}(K)) = N_{tr} cl(M)$. This implies $N_{tr}\Lambda_P cl(f_{N_{tr}}^{-1}(M)) \subseteq f_{N_{tr}}^{-1}(N_{tr}cl(M))$.

(iv) \Rightarrow (iii) Let $M = f_{N_{tr}}(K)$. Then, by assumption, $N_{tr}\Lambda_P cl(K) = N_{tr}\Lambda_P cl\left(f_{N_{tr}}^{-1}(M)\right) \subseteq f_{N_{tr}}^{-1}\left(N_{tr}cl(M)\right) \subseteq f_{N_{tr}}^{-1}\left(N_{tr}cl\left(f_{N_{tr}}(K)\right)\right)$. This implies $f_{N_{tr}}\left(N_{tr}\Lambda_P cl(K)\right) \subseteq N_{tr}cl\left(f_{N_{tr}}(K)\right)$

(iv) \Rightarrow (i) Let M be N_{tr} open in V. Then M^C is N_{tr} closed in V. By assumption, $f_{N_{tr}}^{-1}(M^C) = f_{N_{tr}}^{-1}\left(N_{tr}\,cl(M^C)\right) \supseteq N_{tr}\Lambda_P cl\left(f_{N_{tr}}^{-1}(M^C)\right)$. Also, we know that $f_{N_{tr}}^{-1}(M^C) \subseteq N_{tr}\Lambda_P cl\left(f_{N_{tr}}^{-1}(M^C)\right)$. Hence $f_{N_{tr}}^{-1}(M^C) = N_{tr}\Lambda_P cl\left(f_{N_{tr}}^{-1}(M^C)\right)$. Therefore, $f_{N_{tr}}^{-1}(M^C)$ is $N_{tr}\Lambda_P$ -closed in U. That is, $\left(f_{N_{tr}}^{-1}(M)\right)^C$ is $N_{tr}\Lambda_P$ -closed in U. Hence $f_{N_{tr}}^{-1}(M)$ is $N_{tr}\Lambda_P$ -open in U. Therefore $f_{N_{tr}}$ is $N_{tr}\Lambda_P$ -continuous.

Theorem 4.13: A function $f_{N_{tr}}: (U, \tau_{N_{tr}}) \to (V, \rho_{N_{tr}})$ is $N_{tr} \Lambda_P$ -continuous if and only if $f_{N_{tr}}^{-1}(N_{tr}int(M)) \subseteq N_{tr} \Lambda_Pint(f_{N_{tr}}^{-1}(M))$ for every neutrosophic set M in V.

Proof: Let $f_{N_{tr}}$ be a $N_{tr}\Lambda_P$ -continuous function and M be a neutrosophic set in V. Then $N_{tr}int(M)$ is N_{tr} open in V. By assumption, $f_{N_{tr}}^{-1}(N_{tr}int(M))$ is $N_{tr}\Lambda_P$ -open in U. Now, $f_{N_{tr}}^{-1}(N_{tr}int(M)) \subseteq f_{N_{tr}}^{-1}(M)$ and $N_{tr}\Lambda_Pint\left(f_{N_{tr}}^{-1}(M)\right)$ is the largest $N_{tr}\Lambda_P$ -open set contained in $f_{N_{tr}}^{-1}(M)$. Hence $f_{N_{tr}}^{-1}(N_{tr}int(M)) \subseteq N_{tr}\Lambda_Pint\left(f_{N_{tr}}^{-1}(M)\right)$. Conversely, let M be a N_{tr} open set in V. Then $f_{N_{tr}}^{-1}(M) = f_{N_{tr}}^{-1}(N_{tr}int(M)) \subseteq N_{tr}\Lambda_Pint\left(f_{N_{tr}}^{-1}(M)\right)$. Also, $N_{tr}\Lambda_Pint\left(f_{N_{tr}}^{-1}(M)\right) \subseteq f_{N_{tr}}^{-1}(M)$. This implies $f_{N_{tr}}^{-1}(M)$ is $N_{tr}\Lambda_P$ -open in U. Hence $f_{N_{tr}}$ is $N_{tr}\Lambda_P$ -continuous.

Theorem 4.14: Let $f_{N_{tr}}:(U,\tau_{N_{tr}}) \to (V,\rho_{N_{tr}})$ be a function between two neutrosophic topological spaces. Then the following statements are equivalent:

- i. $f_{N_{tr}}$ is $N_{tr}\Lambda_P$ -continuous.
- ii. For each neutrosophic point $u_{a,b,c}$, the inverse image of every N_{tr} nbhd of $f_{N_{tr}}(u_{a,b,c})$ is $N_{tr}\Lambda_P$ -nbhd of $u_{a,b,c}$.
- iii. For each neutrosophic point $u_{a,b,c}$ in U and every N_{tr} nbhd N of $f_{N_{tr}}(u_{a,b,c})$, there exists a $N_{tr}\Lambda_P$ -open set K in U such that $u_{a,b,c} \in K$ and $f_{N_{tr}}(K) \subseteq N$.

Proof:

(i) \Rightarrow (ii) Let $u_{a,b,c}$ be a neutrosophic point in U and let N be a N_{tr} -nbhd of $f_{N_{tr}}(u_{a,b,c})$. Then there exists a N_{tr} -open set M in V such that $f_{N_{tr}}(u_{a,b,c}) \in M \subseteq N$. Since $f_{N_{tr}}$ is $N_{tr}\Lambda_P$ -continuous, $f_{N_{tr}}^{-1}(M)$ is $N_{tr}\Lambda_P$ -open in U. Also, $u_{a,b,c} \in f_{N_{tr}}^{-1}(f_{N_{tr}}(u_{a,b,c})) \in f_{N_{tr}}^{-1}(M) \subseteq f_{N_{tr}}^{-1}(N)$. Hence there exists a $N_{tr}\Lambda_P$ -open set $f_{N_{tr}}^{-1}(M)$ such that $u_{a,b,c} \in f_{N_{tr}}^{-1}(M) \subseteq f_{N_{tr}}^{-1}(N)$. This implies $f_{N_{tr}}^{-1}(N)$ is $N_{tr}\Lambda_P$ -nbhd of $u_{a,b,c}$.

(ii) \Rightarrow (iii) Let $u_{a,b,c}$ be a neutrosophic point in U and let N be a N_{tr} nbhd of $f_{N_{tr}}(u_{a,b,c})$. Then by assumption, $f_{N_{tr}}^{-1}(N)$ is $N_{tr}\Lambda_P$ -nbhd of $u_{a,b,c}$. Then there exists a $N_{tr}\Lambda_P$ -open set K in U such that $u_{a,b,c} \in K \subseteq f_{N_{tr}}^{-1}(N)$. Thus $u_{a,b,c} \in K$ and $f_{N_{tr}}(K) \subseteq f_{N_{tr}}\left(f_{N_{tr}}^{-1}(N)\right) \subseteq N$.

(iii) \Rightarrow (i) Let M be a N_{tr} open set in V and let $u_{a,b,c} \in f_{N_{tr}}^{-1}(M)$. Since M is N_{tr} open and $f_{N_{tr}}(u_{a,b,c}) \in M$, M is a N_{tr} nbhd of $f_{N_{tr}}(u_{a,b,c})$. Hence it follows (iii) that there exists a $N_{tr}\Lambda_P$ -open set K in U such that $u_{a,b,c} \in K$ and $f_{N_{tr}}(K) \subseteq M$. This implies $u_{a,b,c} \in K \subseteq f_{N_{tr}}^{-1}(f_{N_{tr}}(K)) \subseteq f_{N_{tr}}^{-1}(M)$. By theorem 2.16, $f_{N_{tr}}^{-1}(M)$ is $N_{tr}\Lambda_P$ -open in U. Therefore $f_{N_{tr}}$ is $N_{tr}\Lambda_P$ continuous.





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Definition 4.15: A neutrosophic topological space $(U, \tau_{N_{tr}})$ is said to be $N_{tr}T_{\Lambda_p}$ -space if every $N_{tr}\Lambda_p$ -open set in $(U, \tau_{N_{tr}})$ is N_{tr} open.

Remark 4.16: The composition of two $N_{tr}\Lambda_P$ -continuous functions need not be $N_{tr}\Lambda_P$ -continuous.

Example 4.17: Let $U = \{a, b\}, V = \{x, y\}$ and $W = \{p, q\}$. Consider the neutrosophic topologies $\tau_{N_{tr}} = \{0_{N_{tr}}, K, 1_{N_{tr}}\}$, $\rho_{N_{tr}} = \{0_{N_{tr}}, M, 1_{N_{tr}}\}$ and $\omega_{N_{tr}} = \{0_{N_{tr}}, N, 1_{N_{tr}}\}$ where $K = \{< a, 0.7, 0.5, 0.1 >< b, 0.8, 0.6, 0.2 >\}$, $M = \{< x, 0.2, 0.4, 0.8 >< y, 0.1, 0.5, 0.9 >\}$ and $N = \{< p, 0.2, 0.2, 0.2 >< q, 0.1, 0.30.4 >\}$. Consider the collections $\mathcal{A} = \{A : 0_{N_{tr}} \subset A \subset K\}$, $\mathcal{B} = \{B : K \subset B \subset K^C\}$, $\mathcal{C} = \{C : R \not\subset K; K \not\subset C; C \subset K^C\}$ of neutrosophic sets in U and $\mathcal{D} = \{D : M \subset D \subset 1_{N_{tr}}\}$, the collection of neutrosophic sets in V. Then, $N_{tr}\Lambda_P O(U, \tau_{N_{tr}}) = \{0_{N_{tr}}, K, K^C, \mathcal{A}, \mathcal{B}, \mathcal{C}, 1_{N_{tr}}\}$ and $N_{tr}\Lambda_P O(V, \rho_{N_{tr}}) = \{0_{N_{tr}}, M, \mathcal{D}, 1_{N_{tr}}\}$. Define $f_{N_{tr}}: (U, \tau_{N_{tr}}) \to (V, \rho_{N_{tr}})$ as $f_{N_{tr}}(a) = x$ and $f_{N_{tr}}(b) = y$. Then $f_{N_{tr}}^{-1}(M) = \{< a, 0.2, 0.4, 0.8 >< b, 0.1, 0.5, 0.9 >\}$ is $N_{tr}\Lambda_P$ open in $(U, \tau_{N_{tr}})$. Also, define $g_{N_{tr}}: (V, \rho_{N_{tr}}) \to (W, \omega_{N_{tr}})$ as $g_{N_{tr}}(x) = q$ and $g_{N_{tr}}(y) = p$. Then $g_{N_{tr}}^{-1}(N) = \{< x, 0.1, 0.3, 0.4 >< y, 0.2, 0.2, 0.2, 0.2 >\} \in \mathcal{D}$ which implies $g_{N_{tr}}^{-1}(N)$ is $N_{tr}\Lambda_P$ -open in $(V, \rho_{N_{tr}})$. This implies that both $f_{N_{tr}}$ and $g_{N_{tr}}$ are $N_{tr}\Lambda_P$ -continuous. Now, let $g_{N_{tr}} \circ f_{N_{tr}}: (U, \tau_{N_{tr}}) \to (W, \omega_{N_{tr}})$ be the composition of two $N_{tr}\Lambda_P$ -continuous functions. Then, $g_{N_{tr}} \circ f_{N_{tr}}$ is not $N_{tr}\Lambda_P$ -continuous since $(g_{N_{tr}} \circ f_{N_{tr}})^{-1}(N) = f_{N_{tr}}^{-1}(g_{N_{tr}}^{-1}(N)) = \{< a, 0.1, 0.3, 0.4 >< b, 0.2, 0.2, 0.2, 0.2 >\}$ is not $N_{tr}\Lambda_P$ -open in $(U, \tau_{N_{tr}})$.

Theorem 4.18: Let $(U, \tau_{N_{tr}})$, $(V, \rho_{N_{tr}})$ and $(W, \omega_{N_{tr}})$ be neutrosophic topological space and let $(V, \rho_{N_{tr}})$ be $N_{tr}T_{\Lambda_{P}}$ -space. Then the composition $g_{N_{tr}} \circ f_{N_{tr}} : (U, \tau_{N_{tr}}) \to (W, \omega_{N_{tr}})$ of two $N_{tr}\Lambda_{P}$ -continuous functions $f_{N_{tr}} : (U, \tau_{N_{tr}}) \to (V, \rho_{N_{tr}})$ and $g_{N_{tr}} : (V, \rho_{N_{tr}}) \to (W, \omega_{N_{tr}})$ is $N_{tr}\Lambda_{P}$ -continuous.

Proof: Let N be any N_{tr} open set in W. Since $g_{N_{tr}}$ is $N_{tr}\Lambda_P$ -continuous, $g_{N_{tr}}^{-1}(N)$ is $N_{tr}\Lambda_P$ -open in V. Then, by assumption $g_{N_{tr}}^{-1}(N)$ is N_{tr} open in V. Also, since $f_{N_{tr}}$ is $N_{tr}\Lambda_P$ -continuous, $f_{N_{tr}}^{-1}\left(g_{N_{tr}}^{-1}(N)\right) = \left(g_{N_{tr}} \circ f_{N_{tr}}\right)^{-1}(N)$ is $N_{tr}\Lambda_P$ -open in U. Hence $g_{N_{tr}} \circ f_{N_{tr}}$ is $N_{tr}\Lambda_P$ -continuous.

Theorem 4.19: Let $f_{N_{tr}}: (U, \tau_{N_{tr}}) \to (V, \rho_{N_{tr}})$ be a $N_{tr}\Lambda_P$ -continuous function and $g_{N_{tr}}: (V, \rho_{N_{tr}}) \to (W, \omega_{N_{tr}})$ be a N_{tr} continuous function. Then their composition $g_{N_{tr}} \circ f_{N_{tr}}: (U, \tau_{N_{tr}}) \to (W, \omega_{N_{tr}})$ is $N_{tr}\Lambda_P$ -continuous. **Proof:** Let N be any N_{tr} open set in W. Since $g_{N_{tr}}$ is N_{tr} continuous, $g_{N_{tr}}^{-1}(N)$ is N_{tr} open in V. Also, since $g_{N_{tr}}$ is $N_{tr}\Lambda_P$ -continuous, $f_{N_{tr}}^{-1}(g_{N_{tr}}^{-1}(N)) = (g_{N_{tr}} \circ f_{N_{tr}})^{-1}(N)$ is $N_{tr}\Lambda_P$ -open in V. Hence $g_{N_{tr}} \circ f_{N_{tr}}$ is $N_{tr}\Lambda_P$ -continuous.

Theorem 4.20: Let $f_{N_{tr}}: (U, \tau_{N_{tr}}) \to (V, \rho_{N_{tr}})$ be a $N_{tr}\Lambda_P$ -continuous function where $(U, \tau_{N_{tr}})$ is a $N_{tr}T_{\Lambda_P}$ -space. If C is a subset of U, then the restriction $f_{N_{tr}}|_{C}: (C, \tau_{N_{tr}}^{C}) \to (V, \rho_{N_{tr}})$ is also $N_{tr}\Lambda_P$ -continuous.

Proof: Let M be a N_{tr} open set in V. Since $f_{N_{tr}}$ is $N_{tr}\Lambda_{P}$ -continuous, $f_{N_{tr}}^{-1}(M)$ is $N_{tr}\Lambda_{P}$ -open in U. Now, since U is a $N_{tr}T_{\Lambda_{P}}$ -space, $f_{N_{tr}}^{-1}(M)$ is N_{tr} open in U. Hence $f_{N_{tr}}|_{C}^{-1}(M) = f_{N_{tr}}^{-1}(M) \cap 1_{N_{tr}}^{C}$ is N_{tr} open in C. By theorem 2.14, $f_{N_{tr}}|_{C}^{-1}(M)$ is $N_{tr}\Lambda_{P}$ -open in C. Hence $f_{N_{tr}}|_{C}$ is $N_{tr}\Lambda_{P}$ -continuous.

NEUTROSOPHIC Λ_P-IRRESOLUTE FUNCTIONS

Definition 5.1: Let $(U, \tau_{N_{tr}})$ and $(V, \rho_{N_{tr}})$ be neutrosophic topological spaces. Then the function $f_{N_{tr}}: (U, \tau_{N_{tr}}) \to (V, \rho_{N_{tr}})$ is said to be **neutrosophic**Λ_P-irresolute if $f_{N_{tr}}^{-1}(M)$ is $N_{tr}\Lambda_{P}$ -open in $(U, \tau_{N_{tr}})$ for every $N_{tr}\Lambda_{P}$ -open set M in $(V, \rho_{N_{tr}})$.

Example 5.2:Let $U = \{a, b\}, V = \{x, y\}, \tau_{N_{tr}} = \{0_{N_{tr}}, 1_{N_{tr}}, K_{t}\}$ and $\rho_{N_{tr}} = \{0_{N_{tr}}, 1_{N_{tr}}, M\}$ where $K = \{<\alpha, 0.6, 0.3, 0.5> < b, 0.5, 0.8, 0.4> \}$ and $M = \{<x, 0.5, 0.2, 0.7> < y, 0.2, 0.7, 0.9> \}$. Also, consider the collections $\mathcal{A} = \{A: K \subset A; K^{\mathcal{C}} \subset A\}$ and $\mathcal{B} = \{B: M \subset B; M^{\mathcal{C}} \subset B\}$ of neutrosophic sets in U and V respectively. Then, $N_{tr}\Lambda_{P}O(U, \tau_{N_{tr}}) = \{0_{N_{tr}}, K, \mathcal{A}, 1_{N_{tr}}\}$ and $N_{tr}\Lambda_{P}O(V, \rho_{N_{tr}}) = \{<0_{N_{tr}}, M, \mathcal{B}, 1_{N_{tr}}\}$. Now, let us define $f_{N_{tr}}: (U, \tau_{N_{tr}}) \to (V, \rho_{N_{tr}})$ as $f_{N_{tr}}(a) = x$ and $f_{N_{tr}}(b) = y$. Then, $f_{N_{tr}}^{-1}(M) = \{<\alpha, 0.5, 0.2, 0.7> < b, 0.2, 0.7, 0.9> \} \in \mathcal{A}$ and for each $B \in \mathcal{B}$, there exists some $A \in \mathcal{A}$ such that $f_{N_{tr}}^{-1}(B) = A$. Hence the inverse image of every $N_{tr}\Lambda_{P}$ -open set in V is $N_{tr}\Lambda_{P}$ -open in U. Therefore $f_{N_{tr}}$ is $N_{tr}\Lambda_{P}$ -irresolute.





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Theorem 5.3: Every $N_{tr}\Lambda_P$ -irresolute function is $N_{tr}\Lambda_P$ -continuous.

Proof: Let $f_{N_{tr}}: (U, \tau_{N_{tr}}) \to (V, \rho_{N_{tr}})$ be a $N_{tr} \Lambda_P$ -irresolute function and M be a N_{tr} open set in V. Then, by theorem 2.14, M is $N_{tr} \Lambda_P$ -open in V. Since $f_{N_{tr}}$ is $N_{tr} \Lambda_P$ -irreolsute, $f_{N_{tr}}^{-1}(M)$ is $N_{tr} \Lambda_P$ -open in U. Hence $f_{N_{tr}}$ is $N_{tr} \Lambda_P$ -continuous.

Remark 5.4: The above theorem's converse need not hold.

Theorem 5.6: Let $f_{N_{tr}}: (U, \tau_{N_{tr}}) \to (V, \rho_{N_{tr}})$ be $aN_{tr} \Lambda_P$ -continuous function where $(V, \rho_{N_{tr}})$ is a $N_{tr} T_{\Lambda_P}$ -space. Then $f_{N_{tr}}$ is $N_{tr} \Lambda_P$ -irresolute.

Proof: Let M be $N_{tr}\Lambda_P$ -open in V. Then, by assumption M is N_{tr} open in V. Since $f_{N_{tr}}$ is $N_{tr}\Lambda_P$ -continuous, $f_{N_{tr}}^{-1}(M)$ is $N_{tr}\Lambda_P$ -open in U. Hence $f_{N_{tr}}$ is $N_{tr}\Lambda_P$ -irresolute.

Theorem 5.7: If $f_{N_{tr}}: (U, \tau_{N_{tr}}) \to (V, \rho_{N_{tr}})$ and $g_{N_{tr}}: (V, \rho_{N_{tr}}) \to (W, \omega_{N_{tr}})$ are $N_{tr} \Lambda_P$ -irresolute functions, then their composition $g_{N_{tr}} \circ f_{N_{tr}}: (U, \tau_{N_{tr}}) \to (W, \omega_{N_{tr}})$ is also $N_{tr} \Lambda_P$ -irresolute.

Proof: Let N be $N_{tr}\Lambda_P$ -open in W. Since $g_{N_{tr}}$ is $N_{tr}\Lambda_P$ -irresolute, $g_{N_{tr}}^{-1}(N)$ is $N_{tr}\Lambda_P$ -open in V. Again, since $f_{N_{tr}}$ is $N_{tr}\Lambda_P$ -irresolute, $f_{N_{tr}}^{-1}\left(g_{N_{tr}}^{-1}(N)\right) = \left(g_{N_{tr}} \circ f_{N_{tr}}\right)^{-1}(N)$ is $N_{tr}\Lambda_P$ -open in U. Hence $g_{N_{tr}} \circ f_{N_{tr}}$ is $N_{tr}\Lambda_P$ -irresolute.

Theorem 5.8: If $f_{N_{tr}}: (U, \tau_{N_{tr}}) \to (V, \rho_{N_{tr}})$ is $N_{tr} \Lambda_P$ -irresolute and $g_{N_{tr}}: (V, \rho_{N_{tr}}) \to (W, \omega_{N_{tr}})$ is $N_{tr} \Lambda_P$ -continuous, then $g_{N_{tr}} \circ f_{N_{tr}}: (U, \tau_{N_{tr}}) \to (W, \omega_{N_{tr}})$ is $N_{tr} \Lambda_P$ -continuous.

Proof: Let N be N_{tr} open in W. Since $g_{N_{tr}}$ is $N_{tr}\Lambda_P$ -continuous, $g_{N_{tr}}^{-1}(N)$ is $N_{tr}\Lambda_P$ -open in V. Also, since $f_{N_{tr}}$ is $N_{tr}\Lambda_P$ -irresolute, $f_{N_{tr}}^{-1}\left(g_{N_{tr}}^{-1}(N)\right) = \left(g_{N_{tr}} \circ f_{N_{tr}}\right)^{-1}(N)$ is $N_{tr}\Lambda_P$ -open in U. Hence $g_{N_{tr}} \circ f_{N_{tr}}$ is $N_{tr}\Lambda_P$ -continuous.

Theorem 5.9: Let $f_{N_{tr}}: (U, \tau_{N_{tr}}) \to (V, \rho_{N_{tr}})$ be a function between two neutrosophic topological spaces. Then the following statements are equivalent:

- i. $f_{N_{tr}}$ is $N_{tr} \Lambda_P$ -irresolute.
- ii. The inverse image of every $N_{tr}\Lambda_P$ -closed set in $(V, \rho_{N_{tr}})$ is $N_{tr}\Lambda_P$ -closed in $(U, \tau_{N_{tr}})$.
- iii. $f_{N_{tr}}(N_{tr}\Lambda_P cl(K)) \subseteq N_{tr}\Lambda_P cl(f_{N_{tr}}(K))$ for every neutrosophic set K in U.
- iv. $N_{tr} \Lambda_P cl(f_{N_{tr}}^{-1}(M)) \subseteq f_{N_{tr}}^{-1}(N_{tr} \Lambda_P cl(M))$ for every neutrosophic set M in V.
- v. $f_{N_{tr}}^{-1}(N_{tr}\Lambda_P int(M)) \subseteq N_{tr}\Lambda_P int(f_{N_{tr}}^{-1}(M))$ for every neutrosophic set M in V.

Proof:

(i) \Rightarrow (ii) Let $f_{N_{tr}}$ be a $N_{tr}\Lambda_P$ -irresolute function and M be a $N_{tr}\Lambda_P$ -closed set in V. Then M^c is $N_{tr}\Lambda_P$ -open in V. Since $f_{N_{tr}}$ is $N_{tr}\Lambda_P$ -irresolute, $f_{N_{tr}}^{-1}(M^c)$ is $N_{tr}\Lambda_P$ -open in U. That is, $(f_{N_{tr}}^{-1}(M))^c$ is $N_{tr}\Lambda_P$ -open in U. Hence $f_{N_{tr}}^{-1}(M)$ is $N_{tr}\Lambda_P$ -closed in U.

(ii) \Rightarrow (i) Let M be $N_{tr} \Lambda_P$ -open in V. Then M^c is $N_{tr} \Lambda_P$ -closed in V. By assumption, $f_{N_{tr}}^{-1}(M^c)$ is $N_{tr} \Lambda_P$ -closed in U. That is, $(f_{N_{tr}}^{-1}(M))^c$ is $N_{tr} \Lambda_P$ -closed in U. Hence $f_{N_{tr}}^{-1}(M)$ is $N_{tr} \Lambda_P$ -open in U. Therefore, $f_{N_{tr}}$ is $N_{tr} \Lambda_P$ -irresolute.

(ii) \Rightarrow (iii) Let K be a neutrosophic set in U. Now, $K \subseteq f_{N_{tr}}^{-1}(f_{N_{tr}}(K))$ which implies $K \subseteq f_{N_{tr}}^{-1}(N_{tr}\Lambda_P cl(f_{N_{tr}}(K)))$. Since $N_{tr}\Lambda_P cl(f_{N_{tr}}(K))$ is $N_{tr}\Lambda_P$ -closed in V, by assumption $f_{N_{tr}}^{-1}(N_{tr}\Lambda_P cl(f_{N_{tr}}(K)))$ is a $N_{tr}\Lambda_P$ -closed set containing K.





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Also, $N_{tr}\Lambda_{P}cl(K)$ is the smallest $N_{tr}\Lambda_{P}$ -closed set containing K. Hence, $N_{tr}\Lambda_{P}cl(K) \subseteq f_{N_{tr}}^{-1}\left(N_{tr}\Lambda_{P}cl\left(f_{N_{tr}}(K)\right)\right)$. Therefore, $f_{N_{tr}}\left(N_{tr}\Lambda_{P}cl(K)\right) \subseteq N_{tr}\Lambda_{P}cl\left(f_{N_{tr}}(K)\right)$.

(iii) \Rightarrow (ii) Let M be a $N_{tr}\Lambda_P$ -closed set in V. Then, by assumption $f_{N_{tr}}\left(N_{tr}\Lambda_P cl\left(f_{N_{tr}}^{-1}(M)\right)\right) \subseteq N_{tr}\Lambda_P cl\left(f_{N_{tr}}^{-1}(M)\right) \subseteq N_{tr}\Lambda_P cl\left(f_{N_{tr}}^{-1}(M)\right) \subseteq N_{tr}\Lambda_P cl\left(f_{N_{tr}}^{-1}(M)\right) \subseteq N_{tr}\Lambda_P cl\left(f_{N_{tr}}^{-1}(M)\right) \subseteq I_{N_{tr}}^{-1}(M)$. Also, $f_{N_{tr}}^{-1}(M) \subseteq N_{tr}\Lambda_P cl\left(f_{N_{tr}}^{-1}(M)\right)$. Hence $f_{N_{tr}}^{-1}(M)$ is $N_{tr}\Lambda_P$ -closed in U.

(iii) \Rightarrow (iv) Let M be a neutrosophic set in V and let $K = f_{N_{tr}}^{-1}(M)$. By assumption, $f_{N_{tr}}(N_{tr}\Lambda_{P}cl(K)) \subseteq N_{tr}\Lambda_{P}cl(f_{N_{tr}}(K)) = N_{tr}\Lambda_{P}cl(M)$. This implies $N_{tr}\Lambda_{P}cl(f_{N_{tr}}(M)) \subseteq f_{N_{tr}}^{-1}(N_{tr}\Lambda_{P}cl(M))$.

(iv) \Rightarrow (iii) Let $M = f_{N_{tr}}(K)$. Then, by assumption, $N_{tr}\Lambda_P cl(K) = N_{tr}\Lambda_P cl\left(f_{N_{tr}}^{-1}(M)\right) \subseteq f_{N_{tr}}^{-1}\left(N_{tr}\Lambda_P cl(M)\right) = f_{N_{tr}}^{-1}\left(N_{tr}\Lambda_P cl\left(f_{N_{tr}}(K)\right)\right)$. This implies $f_{N_{tr}}\left(N_{tr}\Lambda_P cl(K)\right) \subseteq N_{tr}\Lambda_P cl\left(f_{N_{tr}}(K)\right)$.

 $(iv) \Leftrightarrow (v)$ This can be proved by taking complements.

(v) \Rightarrow (i) Let M be a $N_{tr}\Lambda_P$ -open set in V. Then $f_{N_{tr}}^{-1}(M) = f_{N_{tr}}^{-1}(N_{tr}\Lambda_P int(M)) \subseteq N_{tr}\Lambda_P int(f_{N_{tr}}^{-1}(M))$. Also, $N_{tr}\Lambda_P int(f_{N_{tr}}^{-1}(M)) \subseteq f_{N_{tr}}^{-1}(M)$. This implies $f_{N_{tr}}^{-1}(M)$ is $N_{tr}\Lambda_P$ -open in U. Hence $f_{N_{tr}}$ is $N_{tr}\Lambda_P$ -irresolute.

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RESEARCH ARTICLE

A New Perspective of Connectedness and Hyperconnectedness in Soft Nano Topological Spaces

P.Anbarasi Rodrigo¹ and P.Subithra^{2*}

¹Assistant Professor, Department of Mathematics, St.Mary's College(Autonomous),(Affiliated to Manonmaniam Sundaranar University, Abishekapatti, Tirunelveli)Thoothukudi, Tamil Nadu, India. ²Research Scholar, Reg.No. 21212212092003, Department of Mathematics, St.Mary's College (Autonomous), (Affiliated to Manonmaniam Sundaranar University, Abishekapatti, Tirunelveli) Thoothukudi, Tamil Nadu, India.

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*Address for Correspondence P.Subithra

Research Scholar,

Reg. No. 21212212092003

Department of Mathematics,

St.Mary's College(Autonomous),

(Affiliated to Manonmaniam Sundaranar University, Abishekapatti, Tirunelveli)

Thoothukudi, Tamil Nadu, India. Email: p.subithra18@gmail.com



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ABSTRACT

The article's primary focus is to introduce soft nano α^* -generalized separated sets and soft nano α^* generalized connected spaces and delve into their characteristics and properties, elucidating their intricate relationship through apt examples. Further, it discuss about the equivalent conditions, union and preservation theorem. Following this examination, the article proceeds to investigate soft nano α^* generalized hyperconnected spaces, particularly through the lens of soft nano α^* -generalized dense sets and explores their correlation with soft nano α^* -generalized connected spaces. Additionally, it presents equivalent conditions, preservation theorems and counterexamples to reverse implications offering a deeper understanding of the complexities involved.

Keywords: soft nano α^* -generalized separated sets, soft nano α^* -generalized connectedness, soft nano α^* generalized dense sets, soft nano α^* -generalized nowhere dense sets, soft nano α^* -generalized hyper connectedness.

MSC Code: Primary 54D05, 54D99, Secondary 54A05





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INTRODUCTION

Molodtsov [10] introduced the soft set theory in 1999 as a mathematical tool to address problems involving imprecise, indeterminacy and inconsistent data, to alleviate challenges in conventional theoretical approaches and proposed the fundamental results of soft set theory. Subsequently, in 2011, Muhammad Shabir and Munazza Naz [11] pioneered the concept of soft topological spaces defining them over an initial universe with a fixed set of parameters. Meanwhile in 2012, the notion of nano topology was unveiled by Lellis Thivagar and Carmel Richard [7], which is an extension of set theory to study the intelligent systems characterized by insufficient and incomplete information and it is defined in terms of approximations and boundary region of a subset of a universe using an equivalence relation on it. By taking inspirations from the above, in 2017, Benchalli et al. [5] introduced the notion of soft nano topological spaces using soft set equivalence relation on the universal set. Connectedness ensures continuity and integrity of spaces, allowing for the study of shapes and relationships critical for understanding complex structures. The notion of connectedness made its advent by the effort of R.L Wilder [14] in 1978. After that, the concept of connectedness was introduced in 2018, by S.Krishnaprakash, R.Ramesh and R.Suresh [6] in nano topology and in 2020, by P.G Patil and S.Benakanawari [12] in soft nano topology. Hyper connectedness explores intricate relationships among points, enabling deep analysis of complex structures and interconnections within spaces. In 1970, L.A. Steen and J.A.Seebach [9] pioneered the concept of hyper connectedness in topological spaces. Subsequently, nano hyperconnected spaces was introduced by M.LellisThivagar and I.Geetha Antoinette [8] in 2019. This paper communicates the role of soft nano α^* -generalized connected space through the lens of soft nano α^* generalized separated sets and also elucidates the notion of soft nano α^* -generalized hyper connected spaces. A new class of sets namely soft nano α^* -generalized separated sets is ideated and exercised with theorems and appropriate examples. Furthermore, novel spaces including soft nano α^* -generalized connected spaces and soft nano α^* generalized hyperconnected spaces are proposed and the essential characteristics, properties and equivalent conditions of these spaces are analysed, providing appropriate exemplifications for counter parts and the preservation theorem is also established.

Preliminaries

In this segment, we have outlined fundamental concepts and findings essential for advancing this work.

Definition 2.1: [3]Let \mathbb{U} be a non-empty finite set of objects called the universe and \mathfrak{W} be the set of parameters. Let \mathcal{X} be a soft equivalence relation on \mathbb{U} . The triplet $(\tau_{\mathcal{R}}\mathcal{X}, \mathbb{U}, \mathfrak{W})$ is said to be the soft approximation space. Let $\mathcal{X} \subseteq_{\mathcal{S}_t \mathcal{N}} \mathbb{U}$. Then

- 1. The **soft lower approximation** of \mathcal{X} with respect to \mathcal{R} and the set of parameters \mathfrak{W} is the set of all objects, which can be for certain classified as \mathcal{X} with respect to \mathcal{R} and it is denoted by $(\mathcal{L}_{\mathcal{R}}(\mathcal{X}), \mathfrak{W})$. That is, $(\mathcal{L}_{\mathcal{R}}(\mathcal{X}), \mathfrak{W}) = \bigcup \{\mathcal{R}(\mathbb{A}) : \mathcal{R}(\mathbb{A}) \subseteq_{\mathcal{S}_r \mathcal{N}} \mathcal{X}\}$, where $\mathcal{R}(\mathbb{A})$ denote the equivalence class determined by $\mathbb{A} \in \mathcal{Y}$.
- 2. The **soft upper approximation** of \mathcal{X} with respect to \mathcal{R} and the set of parameters \mathfrak{W} is the set of all objects, which can be possibly classified as \mathcal{X} with respect to \mathcal{R} and it is denoted by $(\mathcal{U}_{\mathcal{R}}(\mathcal{X}), \mathfrak{W})$. That is, $(\mathcal{U}_{\mathcal{R}}(\mathcal{X}), \mathfrak{W}) = \bigcup \{\mathcal{R}(\mathbb{A}) : \mathcal{R}(\mathbb{A}) \cap \mathcal{X} \neq \emptyset\}$
- 3. The **soft boundary region** of \mathcal{X} with respect to \mathcal{R} and the set of parameters \mathfrak{B} is the set of all objects, which can be classified neither inside \mathcal{X} nor as outside \mathcal{X} with respect to \mathcal{R} and it is denoted by $(\mathcal{B}_{\mathcal{R}}(\mathcal{X}), \mathfrak{B})$. That is, $(\mathcal{B}_{\mathcal{R}}(\mathcal{X}), \mathfrak{B}) = (\mathcal{U}_{\mathcal{R}}(\mathcal{X}), \mathfrak{B}) (\mathcal{L}_{\mathcal{R}}(\mathcal{X}), \mathfrak{B})$.

Definition 2.2: [3]Let \mathbb{U} be a non-empty universal set and \mathfrak{W} be the set of parameters. Let \mathcal{R} be a soft equivalence relation on \mathbb{U} . Let $\mathcal{X} \subseteq_{\mathcal{S}_{\ell}\mathcal{N}} \mathbb{U}$. Let $(\tau_{\mathcal{R}}\mathcal{X}, \mathbb{U}, \mathfrak{W}) = \{\emptyset, (\mathbb{U}, \mathfrak{W}), (\mathcal{L}_{\mathcal{R}}(\mathcal{X}), \mathfrak{W}), (\mathcal{U}_{\mathcal{R}}(\mathcal{X}), \mathfrak{W}), (\mathcal{B}_{\mathcal{R}}(\mathcal{X}), \mathfrak{W})\}$. Then $(\tau_{\mathcal{R}}\mathcal{X}, \mathbb{U}, \mathfrak{W})$ is a Soft topology on $(\mathbb{U}, \mathfrak{W})$, called as the **soft nano topology** with respect to \mathcal{X} . Elements of Soft Nano topology are called **soft nano open sets** $(\mathcal{S}_{t}\mathcal{N} - \mathbf{O}\mathbf{s})$ and $(\tau_{\mathcal{R}}\mathcal{X}, \mathbb{U}, \mathfrak{W})$ is called **soft nano topological space** $(\mathcal{S}_{t}\mathcal{N}T\mathcal{S})$. The complements of $\mathcal{S}_{t}\mathcal{N}\alpha^{*}g - \mathbf{O}\mathbf{s}$ are called **soft nano closed sets** $(\mathcal{S}_{t}\mathcal{N} - \mathbf{C}\mathbf{s})$.

Definition 2.3: [2]A subset $(\mathcal{R}, \mathfrak{B})$ of a $\mathcal{S}_t \mathcal{N} T \mathcal{S}(\tau_{\mathcal{R}} \mathcal{X}, \mathbf{U}, \mathfrak{B})$ is called soft nano α^* -generalized closed $(\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{C} s)$ if $\mathcal{S}_t \mathcal{N} \alpha c l(\mathcal{R}, \mathfrak{B}) \subseteq_{\mathcal{S}_t \mathcal{N}} \mathcal{S}_t \mathcal{N} int^*(\mathcal{S}, \mathfrak{B})$ whenever $(\mathcal{R}, \mathfrak{B}) \subseteq_{\mathcal{S}_t \mathcal{N}} (\mathcal{S}, \mathfrak{B})$ and $(\mathcal{S}, \mathfrak{B}) \in \mathcal{S}_t \mathcal{N} \alpha \mathcal{O}(\tau_{\mathcal{R}} \mathcal{X}, \mathbf{U}, \mathfrak{B})$. The class of all





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 $S_t \mathcal{N} \alpha^* g - Cs$ in $(\tau_{\mathcal{R}} \mathcal{X}, \mathbf{U}, \mathfrak{W})$ is denoted by $S_t \mathcal{N} \alpha^* g C(\tau_{\mathcal{R}} \mathcal{X}, \mathbf{U}, \mathfrak{W})$. The complement of $S_t \mathcal{N} \alpha^* g - Cs$ are called $S_t \mathcal{N} \alpha^* g - \mathbf{0} s$.

Definition 2.4: Let($\tau_{\mathcal{R}}\mathcal{X}$, \mathcal{V} , \mathfrak{W}) and ($\sigma_{\mathcal{R}}\mathcal{Y}$, \mathcal{V} , \mathcal{F}) be two $\mathcal{S}_t\mathcal{N}TS$. Then \mathfrak{f} : ($\tau_{\mathcal{R}}\mathcal{X}$, \mathcal{V} , \mathfrak{W}) \rightarrow ($\sigma_{\mathcal{R}}\mathcal{Y}$, \mathcal{V} , \mathcal{F}) is called

- $S_t \mathcal{N} \alpha^* g$ -continuous[3] if the inverse image of every $S_t \mathcal{N} Cs$ in $(\sigma_{\mathcal{R}} \mathcal{Y}, \mathcal{V}, \mathcal{F})$ is $aS_t \mathcal{N} \alpha^* g Cs$ in $(\tau_{\mathcal{R}} \mathcal{X}, \mathcal{Y}, \mathfrak{Y})$.
- $S_t \mathcal{N} \alpha^* g$ -irresolute[3] if the inverse image of every $S_t \mathcal{N} \alpha^* g Cs$ in $(\sigma_{\mathcal{R}} \mathcal{Y}, \mathcal{V}, \mathcal{F})$ is $aS_t \mathcal{N} \alpha^* g Cs$ in $(\tau_{\mathcal{R}} \mathcal{X}, \mathsf{U}, \mathfrak{B})$.
- $\mathcal{CS}_t \mathcal{N} \alpha^* g$ -continuous[4] if the inverse image of every $\mathcal{S}_t \mathcal{N} Os$ in $(\sigma_{\mathcal{R}} \mathcal{Y}, \mathcal{V}, \mathcal{F})$ is $a\mathcal{S}_t \mathcal{N} \alpha^* g Cs$ in $(\tau_{\mathcal{D}} \mathcal{X}, \mathsf{U}, \mathfrak{B})$.
- $\mathcal{CS}_t \mathcal{N} \alpha^* g$ -irresolute[4] if the inverse image of every $\mathcal{S}_t \mathcal{N} \alpha^* g 0s$ in $(\sigma_{\mathcal{R}} \mathcal{Y}, \mathcal{V}, \mathcal{F})$ is $a\mathcal{S}_t \mathcal{N} \alpha^* g Cs$ in $(\tau_{\mathcal{R}} \mathcal{X}, \mathcal{Y}, \mathcal{Y}, \mathcal{Y})$.

Definition 2.5: [3] A $S_t \mathcal{N} TS(\tau_{\mathcal{R}} \mathcal{X}, \mathcal{Y}, \mathcal{Y}, \mathfrak{B})$ is called $S_t \mathcal{N} \alpha^* g T_{1/2}$ -space if every $S_t \mathcal{N} \alpha^* g - Cs$ of $(\tau_{\mathcal{R}} \mathcal{X}, \mathcal{Y}, \mathcal{Y}, \mathfrak{B})$ is $aS_t \mathcal{N} - Cs$ in $(\tau_{\mathcal{R}} \mathcal{X}, \mathcal{Y}, \mathcal{Y}, \mathfrak{B})$.

Result 2.6: [2]

- a) $S_t \mathcal{N} \alpha^* gint(\mathcal{R}, \mathfrak{W}) = (\mathcal{R}, \mathfrak{W}) \text{ iff } (\mathcal{R}, \mathfrak{W}) \text{ is a } S_t \mathcal{N} \alpha^* g Os \text{ in } (\tau_{\mathcal{R}} \mathcal{X}, \mathcal{U}, \mathfrak{W}).$
- b) $S_t \mathcal{N} \alpha^* gcl(\mathcal{R}, \mathfrak{W}) = (\mathcal{R}, \mathfrak{W}) \text{ iff } (\mathcal{R}, \mathfrak{W}) \text{ is a } S_t \mathcal{N} \alpha^* g \mathcal{C}s \text{ in } (\tau_{\mathcal{R}} \mathcal{X}, \mathbf{U}, \mathfrak{W}).$

Remark 2.7:[2]For any subset $(\mathcal{R}, \mathfrak{W})$ of a $\mathcal{S}_t \mathcal{N}TS$,

- a) $S_t \mathcal{N} \alpha^* gint(\mathcal{R}, \mathfrak{W})^c = (S_t \mathcal{N} \alpha^* gcl(\mathcal{R}, \mathfrak{W}))^c$
- b) $S_t \mathcal{N} \alpha^* gcl(\mathcal{R}, \mathfrak{W})^c = (S_t \mathcal{N} \alpha^* gint(\mathcal{R}, \mathfrak{W}))^c$

Remark 2.8: [3]A function $f:(\tau_{\mathcal{R}}\mathcal{X}, \mathbf{U}, \mathfrak{B}) \to (\sigma_{\mathcal{R}}\mathcal{Y}, \mathcal{V}, \mathcal{F})$ is $\mathcal{S}_t \mathcal{N} \alpha^* g$ –continuous if and only if $f^{-1}(\mathcal{R}, \mathcal{F}) \in \mathcal{S}_t \mathcal{N} \alpha^* g O(\tau_{\mathcal{R}}\mathcal{X}, \mathbf{U}, \mathfrak{B})$ for every $(\mathcal{R}, \mathcal{F}) \in \mathcal{S}_t \mathcal{N} O(\sigma_{\mathcal{R}}\mathcal{Y}, \mathcal{V}, \mathcal{F})$.

Remark 2.9: [3]A function $f: (\tau_{\mathcal{R}} \mathcal{X}, \mathcal{Y}, \mathcal{Y}, \mathcal{Y}) \to (\sigma_{\mathcal{R}} \mathcal{Y}, \mathcal{V}, \mathcal{F})$ is $\mathcal{S}_t \mathcal{N} \alpha^* g$ –irresolute if and only if $\mathcal{S}_t \mathcal{N} \alpha^* gcl(f^{-1}(\mathcal{R}, \mathcal{F})) \subseteq f^{-1}(\mathcal{S}_t \mathcal{N} \alpha^* gcl(\mathcal{R}, \mathcal{F}))$ for each soft nano set $(\mathcal{R}, \mathcal{F})$ in $(\sigma_{\mathcal{R}} \mathcal{Y}, \mathcal{V}, \mathcal{F})$.

Definition 2.10: [12]A $S_t \mathcal{N}TS(\tau_{\mathcal{R}}\mathcal{X}, \mathcal{V}, \mathfrak{D})$ is called a soft nano connected space($S_t \mathcal{N} - C_n S$) if ($\mathcal{V}, \mathfrak{D}$) cannot be represented as the union of two disjoint non-empty $S_t \mathcal{N} - Os$.

Soft nano α^* -generalized separated sets

Separated sets in topology are vital for distinguishing points, characterizing topological properties and finding applications in geometry and analysis. They provide frameword for understanding topological structures and their properties. This segment conceptualizes the idea of soft nano α^* -generalized separated sets in soft nano topology and discuss their properties, validating their counterparts with illustrations and its equivalent conditions.

Definition 3.1: Two non-empty soft nano subsets $(\mathcal{R}, \mathfrak{W})$ and $(\mathcal{S}, \mathfrak{W})$ of a $\mathcal{S}_t \mathcal{N} T S(\tau_{\mathcal{R}} \mathcal{X}, \mathbf{U}, \mathfrak{W})$ are said to be soft nano α^* -generalized separated $(\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{S}_p s)$ if $(\mathcal{R}, \mathfrak{W}) \cap \mathcal{S}_t \mathcal{N} \alpha^* g c l(\mathcal{S}, \mathfrak{W}) = \mathcal{S}_t \mathcal{N} \alpha^* g c l(\mathcal{R}, \mathfrak{W}) \cap (\mathcal{S}, \mathfrak{W}) = \emptyset$.

The following set notations are consistently utilized in the examples of this article.

Let $U = \{i, j, k, \ell\}$ and $\mathfrak{W} = \{m_1, m_2, m_3\}$. Then the soft nano sets are

 $(K_1, \mathfrak{B}) = \{(m_1, \emptyset), (m_2, \emptyset), (m_3, \emptyset)\} = \emptyset$

 $(K_2, \mathfrak{W}) = \{(m_1, \{i\}), (m_2, \{i\}), (m_3, \{i\})\}$

 $(K_3, \mathfrak{B}) = \{(m_1, \{j\}), (m_2, \{j\}), (m_3, \{j\})\}$

 $(K_4, \mathfrak{B}) = \{(m_1, \{k\}), (m_2, \{k\}), (m_3, \{k\})\}$

 $(K_5, \mathfrak{W}) = \{(m_1, \{\ell\}), (m_2, \{\ell\}), (m_3, \{\ell\})\}$





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 \begin{split} & \left( \dot{\mathbf{K}}_{6}, \mathfrak{W} \right) = \left\{ (m_{1}, \{i, j\}), (m_{2}, \{i, j\}), (m_{3}, \{i, j\}) \right\} \\ & \left( \dot{\mathbf{K}}_{7}, \mathfrak{W} \right) = \left\{ (m_{1}, \{i, k\}), (m_{2}, \{i, k\}), (m_{3}, \{i, k\}) \right\} \\ & \left( \dot{\mathbf{K}}_{8}, \mathfrak{W} \right) = \left\{ (m_{1}, \{i, \ell\}), (m_{2}, \{i, \ell\}), (m_{3}, \{i, \ell\}) \right\} \\ & \left( \dot{\mathbf{K}}_{9}, \mathfrak{W} \right) = \left\{ (m_{1}, \{j, k\}), (m_{2}, \{j, k\}), (m_{3}, \{j, k\}) \right\} \\ & \left( \dot{\mathbf{K}}_{10}, \mathfrak{W} \right) = \left\{ (m_{1}, \{j, \ell\}), (m_{2}, \{j, \ell\}), (m_{3}, \{j, \ell\}) \right\} \\ & \left( \dot{\mathbf{K}}_{11}, \mathfrak{W} \right) = \left\{ (m_{1}, \{i, j, k\}), (m_{2}, \{i, j, k\}), (m_{3}, \{i, j, k\}) \right\} \\ & \left( \dot{\mathbf{K}}_{12}, \mathfrak{W} \right) = \left\{ (m_{1}, \{i, j, \ell\}), (m_{2}, \{i, j, \ell\}), (m_{3}, \{i, j, \ell\}) \right\} \\ & \left( \dot{\mathbf{K}}_{13}, \mathfrak{W} \right) = \left\{ (m_{1}, \{i, j, \ell\}), (m_{2}, \{i, j, \ell\}), (m_{3}, \{i, j, \ell\}) \right\} \\ & \left( \dot{\mathbf{K}}_{14}, \mathfrak{W} \right) = \left\{ (m_{1}, \{i, k, \ell\}), (m_{2}, \{i, k, \ell\}), (m_{3}, \{i, k, \ell\}) \right\} \\ & \left( \dot{\mathbf{K}}_{15}, \mathfrak{W} \right) = \left\{ (m_{1}, \{j, k, \ell\}), (m_{2}, \{j, k, \ell\}), (m_{3}, \{j, k, \ell\}) \right\} = (\mathbf{U}, \mathfrak{W}) \end{split}
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Example 3.2: Let $U = \{i, j, k, \ell\}$, $\mathfrak{W} = \{m_1, m_2, m_3\}$ and $\mathcal{X} = \{j, \ell\} \subseteq U$ with $U \setminus \mathcal{R} = \{\{k\}, \{i, j, \ell\}\}$. Then $S_t \mathcal{N} \alpha^* g \mathcal{C}(\tau_{\mathcal{R}} \mathcal{X}, U, \mathfrak{W}) = \{\emptyset, (U, \mathfrak{W}), (K_4, \mathfrak{W}), (K_7, \mathfrak{W}), (K_9, \mathfrak{W}), (K_{11}, \mathfrak{W}), (K_{12}, \mathfrak{W}), (K_{14}, \mathfrak{W}), (K_{15}, \mathfrak{W})\}$. Since $(K_2, \mathfrak{W}) \cap S_t \mathcal{N} \alpha^* g \mathcal{C}(K_5, \mathfrak{W}) = (K_2, \mathfrak{W}) \cap (K_1, \mathfrak{W}) = \emptyset$ and $S_t \mathcal{N} \alpha^* g \mathcal{C}(K_2, \mathfrak{W}) \cap (K_5, \mathfrak{W}) = (K_7, \mathfrak{W}) \cap (K_5, \mathfrak{W}) = \emptyset$, $(K_2, \mathfrak{W}) \cap (K_5, \mathfrak{W}) = S_t \mathcal{N} \alpha^* g - S_v S$.

Theorem 3.3: $S_t \mathcal{N} \alpha^* g - S_{v} s$ are inherently mutually exclusive.

Proof: Assume $(\mathcal{R}, \mathfrak{W})$ and $(\mathcal{S}, \mathfrak{W})$ be two $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{S}_p s$. By definition, $(\mathcal{R}, \mathfrak{W}) \cap \mathcal{S}_t \mathcal{N} \alpha^* g c l(\mathcal{S}, \mathfrak{W}) = \mathcal{S}_t \mathcal{N} \alpha^* g c l(\mathcal{R}, \mathfrak{W}) \cap (\mathcal{S}, \mathfrak{W}) = \emptyset$. Since $(\mathcal{R}, \mathfrak{W}) \cap (\mathcal{S}, \mathfrak{W}) \subseteq_{\mathcal{S}_t \mathcal{N}} (\mathcal{R}, \mathfrak{W}) \cap \mathcal{S}_t \mathcal{N} \alpha^* g c l(\mathcal{S}, \mathfrak{W})$, $(\mathcal{R}, \mathfrak{W}) \cap (\mathcal{S}, \mathfrak{W}) = \emptyset$. Hence $(\mathcal{R}, \mathfrak{W})$ and $(\mathcal{S}, \mathfrak{W})$ are mutually exclusive.

Theorem 3.4: For any two non-empty $S_t \mathcal{N} \alpha^* g - Cs$ that are mutually exclusive, there exists $S_t \mathcal{N} \alpha^* g$ –separation between them.

Proof: Suppose $(\mathcal{R}, \mathfrak{W})$ and $(\mathcal{S}, \mathfrak{W})$ be non-empty $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{C}s$ that are mutually exclusive. Then $(\mathcal{R}, \mathfrak{W}) \cap \mathcal{S}_t \mathcal{N} \alpha^* g c l(\mathcal{S}, \mathfrak{W}) = \mathcal{S}_t \mathcal{N} \alpha^* g c l(\mathcal{R}, \mathfrak{W}) \cap (\mathcal{S}, \mathfrak{W}) = (\mathcal{R}, \mathfrak{W}) \cap (\mathcal{S}, \mathfrak{W}) = \emptyset$. Hence $(\mathcal{R}, \mathfrak{W})$ and $(\mathcal{S}, \mathfrak{W})$ are $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{S}_v s$.

Remark 3.5: The reverse of the aforementioned theorem may not hold true, as demonstrated by the subsequent example.

Example 3.6: Let $U = \{i, j, k, \ell\}$, $\mathfrak{WB} = \{m_1, m_2, m_3\}$ and $\mathcal{X} = \{j, \ell\} \subseteq U$ with $U \setminus \mathcal{R} = \{\{i\}, \{j, k\}, \{\ell\}\}$. Then $\mathcal{S}_t \mathcal{N} \alpha^* g \mathcal{C}(\tau_{\mathcal{R}} \mathcal{X}, U, \mathfrak{W}) = \{\emptyset, (U, \mathfrak{W}), (K_2, \mathfrak{W}), (K_6, \mathfrak{W}), (K_7, \mathfrak{W}), (K_8, \mathfrak{W}), (K_{12}, \mathfrak{W}), (K_{13}, \mathfrak{W}), (K_{14}, \mathfrak{W})\}$. Here $(K_3, \mathfrak{W}) \cap (K_4, \mathfrak{W}) = (K_6, \mathfrak{W}) \cap (K_4, \mathfrak{W}) = (K_6, \mathfrak{W}) \cap (K_4, \mathfrak{W}) = \emptyset$. Hence (K_2, \mathfrak{W}) and (K_5, \mathfrak{W}) are mutually exclusive non-empty $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{S}_p s$ but not $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{S}_s s$.

Theorem 3.7: If $(S_1, \mathfrak{W}), (S_2, \mathfrak{W}) \in S_t \mathcal{N} \alpha^* g \mathcal{O}(\tau_{\mathcal{R}} \mathcal{X}, \centsymbol{U}, \mathfrak{W})$ such that $(\mathcal{R}_1, \mathfrak{W}) = (S_1, \mathfrak{W}) \cap (S_2, \mathfrak{W})^c$ and $(\mathcal{R}_2, \mathfrak{W}) = (S_1, \mathfrak{W})^c \cap (S_2, \mathfrak{W})$, then $(\mathcal{R}_1, \mathfrak{W})$ and $(\mathcal{R}_2, \mathfrak{W})$ are $S_t \mathcal{N} \alpha^* g - S_p s$.

 $\begin{array}{lll} \textbf{Proof:} & \text{Since} & (\mathcal{S}_1,\mathfrak{W}), (\mathcal{S}_2,\mathfrak{W}) \in \mathcal{S}_t \mathcal{N} \alpha^* g \mathcal{O}(\tau_{\mathcal{R}} \mathcal{X}, \cup{U}, \cup{W}), & (\mathcal{S}_1,\mathfrak{W})^c, (\mathcal{S}_2,\mathfrak{W})^c \in \mathcal{S}_t \mathcal{N} \alpha^* g \mathcal{C}(\tau_{\mathcal{R}} \mathcal{X}, \cup{U}, \cup{W}). & \text{Since} \\ (\mathcal{R}_1,\mathfrak{W}) \subseteq_{\mathcal{S}_t \mathcal{N}} (\mathcal{S}_2,\mathfrak{W})^c, & \mathcal{S}_t \mathcal{N} \alpha^* g \mathcal{C}(\mathcal{R}_1,\mathfrak{W}) \subseteq_{\mathcal{S}_t \mathcal{N}} \mathcal{S}_t \mathcal{N} \alpha^* g \mathcal{C}(\mathcal{S}_2,\mathfrak{W})^c = (\mathcal{S}_2,\mathfrak{W})^c. & \text{Thus } \mathcal{S}_t \mathcal{N} \alpha^* g \mathcal{C}(\mathcal{R}_1,\mathfrak{W}) \cap (\mathcal{S}_2,\mathfrak{W}) = \emptyset \\ \text{which implies} & \mathcal{S}_t \mathcal{N} \alpha^* g \mathcal{C}(\mathcal{R}_1,\mathfrak{W}) \cap (\mathcal{R}_2,\mathfrak{W}) = \emptyset, & \text{since} & (\mathcal{R}_2,\mathfrak{W}) \subseteq_{\mathcal{S}_t \mathcal{N}} (\mathcal{S}_2,\mathfrak{W}). & \text{Similarly,} & \text{by taking} \\ (\mathcal{R}_2,\mathfrak{W}) \subseteq_{\mathcal{S}_t \mathcal{N}} (\mathcal{S}_1,\mathfrak{W})^c, & \mathcal{S}_t \mathcal{N} \alpha^* g \mathcal{C}(\mathcal{R}_1,\mathfrak{W}) \cap \mathcal{S}_t \mathcal{N} \alpha^* g \mathcal{C}(\mathcal{R}_2,\mathfrak{W}) = \emptyset. & \text{Hence } (\mathcal{R}_1,\mathfrak{W}) & \text{and } (\mathcal{R}_2,\mathfrak{W}) & \text{are } \mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{S}_p \mathcal{S}. \\ \end{array}$

Theorem 3.8: If $(\mathcal{R}_2, \mathfrak{W})$ and $(\mathcal{S}_2, \mathfrak{W})$ are $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{S}_p s$ such that $(\mathcal{R}_1, \mathfrak{W}) \subseteq_{\mathcal{S}_t \mathcal{N}} (\mathcal{R}_2, \mathfrak{W})$ and $(\mathcal{S}_1, \mathfrak{W}) \subseteq_{\mathcal{S}_t \mathcal{N}} (\mathcal{S}_2, \mathfrak{W})$, then $(\mathcal{R}_1, \mathfrak{W})$ and $(\mathcal{S}_1, \mathfrak{W})$ are $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{S}_p s$.

Proof: Let $(\mathcal{R}_2, \mathfrak{B})$ and $(\mathcal{S}_2, \mathfrak{B})$ are $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{S}_{p} s$. Then $(\mathcal{R}_2, \mathfrak{B}) \cap \mathcal{S}_t \mathcal{N} \alpha^* g c l(\mathcal{S}_2, \mathfrak{B}) = \mathcal{S}_t \mathcal{N} \alpha^* g c l(\mathcal{R}_2, \mathfrak{B}) \cap (\mathcal{S}_2, \mathfrak{B}) = \emptyset$. Since $(\mathcal{R}_1, \mathfrak{B}) \subseteq_{\mathcal{S}_t \mathcal{N}} (\mathcal{R}_2, \mathfrak{B})$ and $(\mathcal{S}_1, \mathfrak{B}) \subseteq_{\mathcal{S}_t \mathcal{N}} (\mathcal{S}_2, \mathfrak{B})$, $(\mathcal{R}_2, \mathfrak{B}) \cap \mathcal{S}_t \mathcal{N} \alpha^* g c l(\mathcal{S}_1, \mathfrak{B}) \subseteq_{\mathcal{S}_t \mathcal{N}} (\mathcal{R}_2, \mathfrak{B}) \cap \mathcal{S}_t \mathcal{N} \alpha^* g c l(\mathcal{S}_2, \mathfrak{B}) = \emptyset$.





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 \emptyset and $\mathcal{S}_t \mathcal{N} \alpha^* gcl(\mathcal{R}_1, \mathfrak{W}) \cap (\mathcal{S}_1, \mathfrak{W}) \subseteq_{\mathcal{S}_t \mathcal{N}} \mathcal{S}_t \mathcal{N} \alpha^* gcl(\mathcal{R}_2, \mathfrak{W}) \cap (\mathcal{S}_2, \mathfrak{W}) = \emptyset$. Hence $(\mathcal{R}_1, \mathfrak{W})$ and $(\mathcal{S}_1, \mathfrak{W})$ are $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{S}_v \mathcal{S}$.

Theorem 3.9: The soft nano subsets $(\mathcal{R}_1, \mathfrak{B})$ and $(\mathcal{S}_1, \mathfrak{B})$ of a $\mathcal{S}_t \mathcal{N} T S(\tau_{\mathcal{R}} \mathcal{X}, \mathbf{U}, \mathfrak{B})$ are $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{S}_{\mathcal{P}} s$ if and only if there exists $\mathcal{S}_t \mathcal{N} \alpha^* g - Os(\mathcal{R}_2, \mathfrak{B})$ and $(\mathcal{S}_2, \mathfrak{B})$ such that $(\mathcal{R}_1, \mathfrak{B}) \subseteq_{\mathcal{S}_t \mathcal{N}} (\mathcal{R}_2, \mathfrak{B})$ and $(\mathcal{S}_1, \mathfrak{B}) \subseteq_{\mathcal{S}_t \mathcal{N}} (\mathcal{S}_2, \mathfrak{B})$, and $(\mathcal{S}_1, \mathfrak{B}) \subseteq_{\mathcal{S}_t \mathcal{N}} (\mathcal{S}_2, \mathfrak{B})$ and $(\mathcal{S}_1, \mathfrak{B}) \cap (\mathcal{R}_2, \mathfrak{B}) = \emptyset$ and $(\mathcal{S}_1, \mathfrak{B}) \cap (\mathcal{R}_2, \mathfrak{B}) = \emptyset$.

Proof: Assume $(\mathcal{R}_1, \mathfrak{W})$ and $(\mathcal{S}_1, \mathfrak{W})$ are $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{S}_p s$. Then $(\mathcal{R}_1, \mathfrak{W}) \cap \mathcal{S}_t \mathcal{N} \alpha^* g c l(\mathcal{S}_1, \mathfrak{W}) = \mathcal{S}_t \mathcal{N} \alpha^* g c l(\mathcal{R}_1, \mathfrak{W}) \cap \mathcal{S}_t \mathcal{N} \alpha^* g c l(\mathcal{S}_1, \mathfrak{$

Conversely, assume that $(\mathcal{R}_2, \mathfrak{B}), (\mathcal{S}_2, \mathfrak{B}) \in \mathcal{S}_t \mathcal{N} \alpha^* g O(\tau_{\mathcal{R}} \mathcal{X}, \cup{U}, \mathfrak{B})$ such that $(\mathcal{R}_1, \mathfrak{B}) \subseteq_{\mathcal{S}_t \mathcal{N}} (\mathcal{R}_2, \mathfrak{B})$ and $(\mathcal{S}_1, \mathfrak{B}) \subseteq_{\mathcal{S}_t \mathcal{N}} (\mathcal{S}_2, \mathfrak{B}), (\mathcal{R}_1, \mathfrak{B}) \cap (\mathcal{S}_2, \mathfrak{B}) = \emptyset$ and $(\mathcal{S}_1, \mathfrak{B}) \cap (\mathcal{R}_2, \mathfrak{B}) = \emptyset$. Then $(\mathcal{R}_1, \mathfrak{B}) \subseteq_{\mathcal{S}_t \mathcal{N}} (\cup{U}, \mathfrak{B}) \setminus (\mathcal{S}_2, \mathfrak{B})$ and $(\mathcal{S}_1, \mathfrak{B}) \subseteq_{\mathcal{S}_t \mathcal{N}} (\cup{U}, \mathfrak{B}) \setminus (\mathcal{R}_2, \mathfrak{B}), (\cup{U}, \mathfrak{B}) \setminus (\mathcal{S}_2, \mathfrak{B}) \in \mathcal{S}_t \mathcal{N} \alpha^* g C(\tau_{\mathcal{R}} \mathcal{X}, \cup{U}, \mathfrak{B}).$ Then $\mathcal{S}_t \mathcal{N} \alpha^* g c l(\mathcal{R}_1, \mathfrak{B}) \subseteq_{\mathcal{S}_t \mathcal{N}} (\cup{U}, \mathfrak{B}) \setminus (\mathcal{S}_2, \mathfrak{B})$ and $(\mathcal{S}_1, \mathfrak{B}) \subseteq_{\mathcal{S}_t \mathcal{N}} (\cup{U}, \mathfrak{B}) \setminus (\mathcal{S}_2, \mathfrak{B})$. Since $(\mathcal{R}_1, \mathfrak{B}) \subseteq_{\mathcal{S}_t \mathcal{N}} (\mathcal{R}_2, \mathfrak{B})$ and $(\mathcal{S}_1, \mathfrak{B}) \subseteq_{\mathcal{S}_t \mathcal{N}} (\mathcal{S}_2, \mathfrak{B}), \mathcal{S}_t \mathcal{N} \alpha^* g c l(\mathcal{R}_1, \mathfrak{B}) \subseteq_{\mathcal{S}_t \mathcal{N}} (\cup{U}, \mathfrak{B}) \setminus (\mathcal{S}_1, \mathfrak{B})$ and $(\mathcal{S}_1, \mathfrak{B}) \cap (\mathcal{S}_1, \mathfrak{B}) \cap (\mathcal$

Soft nano α^* -generalized connectedness

Connectedness is a fundamental concept in topology, defining the continuity of a topological space. This property captures the idea of a single, cohesive entity, crucial for understanding the structure and behaviour of topological spaces in various mathematical and real world contexts. This segment underpins the origination of soft nano α^* -generalized connectedness, analyzing the structure and properties, making it a cornerstone of the field.

Definition 4.1: A $S_t \mathcal{N} TS(\tau_{\mathcal{R}} \mathcal{X}, \c{U}, \cents)$ is called soft nano α^* -generalized connected space $(S_t \mathcal{N} \alpha^* g - \mathcal{C}_n S)$ if (\c{U}, \c{W}) cannot be represented as the combination of two non-empty mutually exclusive $S_t \mathcal{N} \alpha^* g - Os$ of $(\tau_{\mathcal{R}} \mathcal{X}, \c{U}, \cal{W})$. A subset of $(\tau_{\mathcal{R}} \mathcal{X}, \c{U}, \cal{W})$ is $S_t \mathcal{N} \alpha^* g - \mathcal{C}_n$ if it is $S_t \mathcal{N} \alpha^* g - \mathcal{C}_n$ as a subspace.

Remark 4.3: A subset of $(\tau_{\mathcal{R}}\mathcal{X}, \mathbf{U}, \mathfrak{W})$ is $\mathcal{S}_t \mathcal{N} \alpha^* g$ –disconnected $(\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{D} \mathcal{C}_n)$ if and only if it is not $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{C}_n$.

Theorem 4.4: For any $S_t \mathcal{N}TS(\tau_{\mathcal{R}} \mathcal{X}, \mathbf{U}, \mathfrak{B})$, the subsequent assertions are equivalent.

- i) $(\tau_{\mathcal{R}} \mathcal{X}, \mathcal{Y}, \mathfrak{Y})$ is a $\mathcal{S}_t \mathcal{N} \alpha^* g \mathcal{C}_n \mathcal{S}$.
- ii) (U, \mathfrak{W}) and \emptyset are the only subsets of (U, \mathfrak{W}) which are both $\mathcal{S}_t \mathcal{N} \alpha^* g Os$ and $\mathcal{S}_t \mathcal{N} \alpha^* g Cs$ in $(\tau_{\mathcal{R}} \mathcal{X}, U, \mathfrak{W})$.
- iii) Each $S_t \mathcal{N} \alpha^* g$ –continuous function of $(\tau_{\mathcal{R}} \mathcal{X}, \mathbf{U}, \mathfrak{W})$ into a $S_t \mathcal{N}$ –discrete space $(\sigma_{\mathcal{R}} \mathcal{Y}, \mathcal{V}, H)$ with at least two points is a constant function.

Proof: i) \Rightarrow ii) Assume($\mathcal{R}, \mathfrak{W}$) $\in \mathcal{S}_t \mathcal{N} \alpha^* g \mathcal{C} \mathcal{O}(\tau_{\mathcal{R}} \mathcal{X}, \underline{\mathbb{U}}, \mathfrak{W})$. Then $(\mathcal{R}, \mathfrak{W})^c \in \mathcal{S}_t \mathcal{N} \alpha^* g \mathcal{C} \mathcal{O}(\tau_{\mathcal{R}} \mathcal{X}, \underline{\mathbb{U}}, \mathfrak{W})$. Thus $(\underline{\mathbb{U}}, \mathfrak{W}) = (\mathcal{R}, \mathfrak{W}) \cup (\mathcal{R}, \mathfrak{W})^c$ and $\emptyset = (\mathcal{R}, \mathfrak{W}) \cap (\mathcal{R}, \mathfrak{W})^c$, where $(\mathcal{R}, \mathfrak{W}), (\mathcal{R}, \mathfrak{W})^c \in \mathcal{S}_t \mathcal{N} \alpha^* g \mathcal{O}(\tau_{\mathcal{R}} \mathcal{X}, \underline{\mathbb{U}}, \mathfrak{W})$. Since $(\tau_{\mathcal{R}} \mathcal{X}, \underline{\mathbb{U}}, \mathfrak{W})$ is a $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{C}_n \mathcal{S}$, either $(\mathcal{R}, \mathfrak{W}) = \emptyset$ or $(\mathcal{R}, \mathfrak{W}) = (\underline{\mathbb{U}}, \mathfrak{W})$.

ii) \Longrightarrow i) Suppose $(\tau_{\mathcal{R}}\mathcal{X}, \Dot{U}, \Bar{W})$ is $\mathcal{S}_t\mathcal{N}\alpha^*g - \mathcal{D}\mathcal{C}_n$. Then $(\Dot{U}, \Bar{W}) = (\mathcal{R}, \Bar{W}) \cup (\mathcal{S}, \Bar{W})$ where (\mathcal{R}, \Bar{W}) and (\mathcal{S}, \Bar{W}) are non-empty mutually exclusive $\mathcal{S}_t\mathcal{N}\alpha^*g - \mathcal{O}s$. Since $(\mathcal{R}, \Bar{W}) = (\Dot{U}, \Bar{W}) \setminus (\mathcal{S}, \Bar{W})$, (\mathcal{R}, \Bar{W}) is $\mathcal{S}_t\mathcal{N}\alpha^*g - \mathcal{C}s$. Since $(\mathcal{R}, \Bar{W}) \in \mathcal{S}_t\mathcal{N}\alpha^*g\mathcal{C}O(\tau_{\mathcal{R}}\mathcal{X}, \Dot{U}, \Bar{W})$, by our assumption, either $(\mathcal{R}, \Bar{W}) = \emptyset$ or $(\mathcal{R}, \Bar{W}) = (\Dot{U}, \Bar{W})$ which implies either $(\mathcal{R}, \Bar{W}) = \emptyset$ or $(\mathcal{S}, \Bar{W}) = \emptyset$, leading to a contradiction. Hence $(\tau_{\mathcal{R}}\mathcal{X}, \Dot{U}, \Bar{W})$ is a $\mathcal{S}_t\mathcal{N}\alpha^*g - \mathcal{C}_n\mathcal{S}$.





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ii) \Longrightarrow iii) Let $\mathfrak{f}: (\tau_{\mathcal{R}} \mathcal{X}, \c U, \mathfrak{M}) \to (\sigma_{\mathcal{R}} \mathcal{Y}, \mathcal{V}, H)$ be $\mathcal{S}_t \mathcal{N} \alpha^* g$ —continuous such that $(\sigma_{\mathcal{R}} \mathcal{Y}, \mathcal{V}, H)$ is a $\mathcal{S}_t \mathcal{N}$ —discrete space with at least two points. Then for each $(\mathcal{R}, H) \in (\sigma_{\mathcal{R}} \mathcal{Y}, \mathcal{V}, H)$, $(\mathcal{R}, H) \in \mathcal{S}_t \mathcal{N} \mathcal{C} O(\sigma_{\mathcal{R}} \mathcal{Y}, \mathcal{V}, H)$. Since \mathfrak{f} is $\mathcal{S}_t \mathcal{N} \alpha^* g$ —continuous, $\mathfrak{f}^{-1}(\mathcal{R}, H) \in \mathcal{S}_t \mathcal{N} \alpha^* g \mathcal{C} O(\tau_{\mathcal{R}} \mathcal{X}, \c U, \c M)$. Hence $(\tau_{\mathcal{R}} \mathcal{X}, \c U, \c M)$ is covered by $\mathcal{S}_t \mathcal{N} \alpha^* g$ —clopen set $\{\mathfrak{f}^{-1}(\mathcal{R}, H) : (\mathcal{R}, H) \in (\sigma_{\mathcal{R}} \mathcal{Y}, \mathcal{V}, H)\}$. Then by our assumption, $\mathfrak{f}^{-1}(\mathcal{R}, H) = \emptyset$ or $(\c U, \c M)$. If $\mathfrak{f}^{-1}(\mathcal{R}, H) = \emptyset$ for all $(\c R, H) \in (\sigma_{\mathcal{R}} \mathcal{Y}, \mathcal{V}, H)$, then \mathfrak{f} fails to be a function. Hence, there exists only one point $(\c R, H) \in (\sigma_{\mathcal{R}} \mathcal{Y}, \mathcal{V}, H)$ such that $\mathfrak{f}^{-1}(\mathcal{R}, H) = (\c U, \c M)$ which shows that \mathfrak{f} is a constant function.

iii) \Rightarrow ii) Let($\mathcal{R}, \mathfrak{W}$) $\in \mathcal{S}_t \mathcal{N} \alpha^* g \mathcal{C} O(\tau_{\mathcal{R}} \mathcal{X}, \cU, \mathfrak{W})$. Suppose $(\mathcal{R}, \mathfrak{W}) \neq \emptyset$. Define $f: (\tau_{\mathcal{R}} \mathcal{X}, \cU, \mathfrak{W}) \to (\sigma_{\mathcal{R}} \mathcal{Y}, \cV, H)$ where $(\sigma_{\mathcal{R}} \mathcal{Y}, \cV, H)$ is a $\mathcal{S}_t \mathcal{N}$ -discrete space with at least two points by $f(\mathcal{R}, \mathfrak{W}) = (\mathcal{S}, H), f((\cU, \mathfrak{W}) \setminus (\mathcal{R}, \mathfrak{W})) = (\mathcal{T}, H)$. Then f is $\mathcal{S}_t \mathcal{N} \alpha^* g$ -continuous. Therefore by hypothesis, f is a constant function. Thus $(\mathcal{S}, H) = (\mathcal{T}, H)$ and hence $(\mathcal{R}, \mathfrak{W}) = (\cU, \mathfrak{W})$.

Theorem 4.5: Every $S_t \mathcal{N} \alpha^* g - C_n S$ is $S_t \mathcal{N} - C_n$.

Proof: Suppose $(\tau_{\mathcal{R}}\mathcal{X}, \underline{\mathbb{U}}, \mathfrak{W})$ is a $\mathcal{S}_t\mathcal{N}\alpha^*g - \mathcal{C}_n\mathcal{S}$ but not $\mathcal{S}_t\mathcal{N} - \mathcal{C}_n$. Then there exists mutually exclusive non-empty $\mathcal{S}_t\mathcal{N} - \mathcal{O}_s(\mathcal{R}, \mathfrak{W})$ and $(\mathcal{S}, \mathfrak{W})$ such that $(\underline{\mathbb{U}}, \mathfrak{W}) = (\mathcal{R}, \mathfrak{W}) \cup (\mathcal{S}, \mathfrak{W})$. Since every $\mathcal{S}_t\mathcal{N} - \mathcal{O}_s$ is a $\mathcal{S}_t\mathcal{N}\alpha^*g - \mathcal{O}_s$, $(\mathcal{R}, \mathfrak{W})$ and $(\mathcal{S}, \mathfrak{W})$ are mutually exclusive non-empty $\mathcal{S}_t\mathcal{N}\alpha^*g - \mathcal{O}_s$ such that $(\underline{\mathbb{U}}, \mathfrak{W}) = (\mathcal{R}, \mathfrak{W}) \cup (\mathcal{S}, \mathfrak{W})$. Then the $\mathcal{S}_t\mathcal{N}\alpha^*g$ –disconnection of $(\tau_{\mathcal{R}}\mathcal{X}, \underline{\mathbb{U}}, \mathfrak{W})$ presents a contradiction. Hence $(\tau_{\mathcal{R}}\mathcal{X}, \underline{\mathbb{U}}, \mathfrak{W})$ is $\mathcal{S}_t\mathcal{N} - \mathcal{C}_n$.

Remark 4.6: Every $S_t \mathcal{N} - C_n S$ is $S_t \mathcal{N} \alpha^* g - C_n$ only if it is a $S_t \mathcal{N} \alpha^* g T_{1/2}$ –space.

Theorem 4.7: For any $\mathcal{S}_t \mathcal{N}TS(\tau_{\mathcal{R}} \mathcal{X}, U, \mathfrak{W})$, the subsequent assertions are equivalent.

- i) $(\tau_{\mathcal{R}} \mathcal{X}, \mathcal{Y}, \mathfrak{W})$ is a $\mathcal{S}_t \mathcal{N} \alpha^* g \mathcal{C}_n S$.
- ii) (U, \mathfrak{B}) cannot be represented as the combination of two mutually exclusive non-empty $S_t \mathcal{N} \alpha^* g Cs$.

Proof: i) \Rightarrow ii) Suppose(\S , \mathfrak{B}) = (\S , \mathfrak{B}) \cup (\S , \mathfrak{B}) where (\S , \mathfrak{B}) and (\S , \mathfrak{B}) are two mutually exclusive non-empty $S_t \mathcal{N} \alpha^* g - Cs$. Since (\S , \mathfrak{B}) = (\S , \mathfrak{B}) (\S , \mathfrak{B}) and (\S , \mathfrak{B}) = (\S , \mathfrak{B}), (\S , \mathfrak{B}), (\S , \mathfrak{B}), (\S , \mathfrak{B}), (\S , \mathfrak{B}) is $S_t \mathcal{N} \alpha^* g - \mathcal{D} \mathcal{C}_n$, which leads to a contradiction of our assumption.

ii) \Rightarrow i) Suppose($\tau_{\mathcal{R}}\mathcal{X}, \mathbf{U}, \mathfrak{W}$) is $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{D} \mathcal{C}_n$. Then $(\mathbf{U}, \mathfrak{W}) = (\mathcal{R}, \mathfrak{W}) \cup (\mathcal{S}, \mathfrak{W})$ where $(\mathcal{R}, \mathfrak{W})$ and $(\mathcal{S}, \mathfrak{W})$ are two mutually exclusive non-empty $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{O} s$. Since $(\mathcal{R}, \mathfrak{W}) = (\mathbf{U}, \mathfrak{W}) \setminus (\mathcal{S}, \mathfrak{W})$ and $(\mathcal{S}, \mathfrak{W}) = (\mathbf{U}, \mathfrak{W}) \setminus (\mathcal{R}, \mathfrak{W})$, $(\mathcal{R}, \mathfrak{W}), (\mathcal{S}, \mathfrak{W}) \in \mathcal{S}_t \mathcal{N} \alpha^* g \mathcal{C}(\tau_{\mathcal{R}} \mathcal{X}, \mathbf{U}, \mathfrak{W})$, which leads to a contradiction of our assumption.

Theorem 4.8:

- i) If $f: (\tau_{\mathcal{R}} \mathcal{X}, \mathbf{U}, \mathfrak{B}) \to (\sigma_{\mathcal{R}} \mathcal{Y}, \mathcal{V}, H)$ is a $\mathcal{S}_t \mathcal{N} \alpha^* g$ —continuous surjection and $(\tau_{\mathcal{R}} \mathcal{X}, \mathbf{U}, \mathfrak{B})$ is $\mathcal{S}_t \mathcal{N} \alpha^* g \mathcal{C}_n$, then $(\sigma_{\mathcal{R}} \mathcal{Y}, \mathcal{V}, H)$ is $\mathcal{S}_t \mathcal{N} \mathcal{C}_n$.
- ii) If $f: (\tau_{\mathcal{R}} \mathcal{X}, \mathbf{U}, \mathfrak{B}) \to (\sigma_{\mathcal{R}} \mathcal{Y}, \mathcal{V}, H)$ is a $\mathcal{S}_t \mathcal{N} \alpha^* g$ –irresolute surjection and $(\tau_{\mathcal{R}} \mathcal{X}, \mathbf{U}, \mathfrak{B})$ is $\mathcal{S}_t \mathcal{N} \alpha^* g \mathcal{C}_n$, then $(\sigma_{\mathcal{R}} \mathcal{Y}, \mathcal{V}, H)$ is $\mathcal{S}_t \mathcal{N} \alpha^* g \mathcal{C}_n$.
- iii) If $f: (\tau_{\mathcal{R}} \mathcal{X}, \mathbf{U}, \mathfrak{B}) \to (\sigma_{\mathcal{R}} \mathcal{Y}, \mathcal{V}, H)$ is a $\mathcal{CS}_t \mathcal{N} \alpha^* g$ –continuous surjection and $(\tau_{\mathcal{R}} \mathcal{X}, \mathbf{U}, \mathfrak{B})$ is $\mathcal{S}_t \mathcal{N} \alpha^* g \mathcal{C}_n$, then $(\sigma_{\mathcal{R}} \mathcal{Y}, \mathcal{V}, H)$ is $\mathcal{S}_t \mathcal{N} \mathcal{C}_n$.
- iv) If $f: (\tau_{\mathcal{R}} \mathcal{X}, \mathbf{U}, \mathfrak{B}) \to (\sigma_{\mathcal{R}} \mathcal{Y}, \mathcal{V}, H)$ is a $\mathcal{CS}_t \mathcal{N} \alpha^* g$ –irresolute surjection and $(\tau_{\mathcal{R}} \mathcal{X}, \mathbf{U}, \mathfrak{B})$ is $\mathcal{S}_t \mathcal{N} \alpha^* g \mathcal{C}_n$, then $(\sigma_{\mathcal{R}} \mathcal{Y}, \mathcal{V}, H)$ is $\mathcal{S}_t \mathcal{N} \alpha^* g \mathcal{C}_n$.

Proof: i) Suppose $(\sigma_{\mathcal{R}}\mathcal{Y}, \mathcal{V}, H)$ is not a $\mathcal{S}_t \mathcal{N} - \mathcal{C}_n \mathcal{S}$. Then $(\mathcal{V}, H) = (\mathcal{R}, H) \cup (\mathcal{S}, H)$ such that (\mathcal{R}, H) and (\mathcal{S}, H) are mutually exclusive non-empty $\mathcal{S}_t \mathcal{N} - \mathcal{O}s$ in $(\sigma_{\mathcal{R}}\mathcal{Y}, \mathcal{V}, H)$. Since f is $\mathcal{S}_t \mathcal{N} \alpha^* g$ –continuous surjection, $(\mathcal{V}, \mathcal{W}) = f^{-1}(\mathcal{V}, H) = f^{-1}(\mathcal{R}, H) \cup (\mathcal{S}, H) = f^{-1}(\mathcal{R}, H) \cup f^{-1}(\mathcal{S}, H)$, where $f^{-1}(\mathcal{R}, H)$ and $f^{-1}(\mathcal{S}, H)$ are mutually exclusive non-empty $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{O}s$ in $(\tau_{\mathcal{R}} \mathcal{X}, \mathcal{V}, \mathcal{W})$. This contradicts the fact that $(\tau_{\mathcal{R}} \mathcal{X}, \mathcal{V}, \mathcal{W})$ is $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{C}_n$. Hence $(\sigma_{\mathcal{R}} \mathcal{Y}, \mathcal{V}, H)$ is $\mathcal{S}_t \mathcal{N} - \mathcal{C}_n$.

Proofs of ii), iii) and iv) are similar to i).

Theorem 4.9: A $S_t \mathcal{N}TS(\tau_{\mathcal{R}} \mathcal{X}, U, \mathfrak{B})$ is a $S_t \mathcal{N} \alpha^* g - C_n S$ if and only if (U, \mathfrak{B}) cannot be represented as a combination of any two $S_t \mathcal{N} \alpha^* g - S_p S$.

Proof: Assume $(\tau_{\mathcal{R}} \mathcal{X}, \mathbf{U}, \mathfrak{B})$ is a $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{C}_n S$. Suppose $(\mathbf{U}, \mathfrak{B}) = (\mathcal{R}, \mathfrak{B}) \cup (\mathcal{S}, \mathfrak{B})$ where $(\mathcal{R}, \mathfrak{B})$ and $(\mathcal{S}, \mathfrak{B})$ are $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{S}_{\mathcal{P}} S$. Then $(\mathcal{R}, \mathfrak{B}) \cap \mathcal{S}_t \mathcal{N} \alpha^* g c l(\mathcal{S}, \mathfrak{B}) = \emptyset$ and $\mathcal{S}_t \mathcal{N} \alpha^* g c l(\mathcal{R}, \mathfrak{B}) \cap (\mathcal{S}, \mathfrak{B}) = \emptyset$, which implies





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 $S_t \mathcal{N} \alpha^* gcl(\mathcal{S}, \mathfrak{W}) \subseteq_{\mathcal{S}_t \mathcal{N}} (\underline{\mathbb{U}}, \mathfrak{W}) \setminus (\mathcal{R}, \mathfrak{W}) = (\mathcal{S}, \mathfrak{W})$ and $S_t \mathcal{N} \alpha^* gcl(\mathcal{R}, \mathfrak{W}) \subseteq_{\mathcal{S}_t \mathcal{N}} (\underline{\mathbb{U}}, \mathfrak{W}) \setminus (\mathcal{S}, \mathfrak{W}) = (\mathcal{R}, \mathfrak{W})$. Thus $(\mathcal{R}, \mathfrak{W}), (\mathcal{S}, \mathfrak{W}) \in S_t \mathcal{N} \alpha^* gC(\tau_{\mathcal{R}} \mathcal{X}, \underline{\mathbb{U}}, \mathfrak{W})$. Since $(\underline{\mathbb{U}}, \mathfrak{W}) = (\mathcal{R}, \mathfrak{W}) \cup (\mathcal{S}, \mathfrak{W}), (\mathcal{R}, \mathfrak{W}), (\mathcal{S}, \mathfrak{W}) \in S_t \mathcal{N} \alpha^* gO(\tau_{\mathcal{R}} \mathcal{X}, \underline{\mathbb{U}}, \mathfrak{W})$. Since $(\mathcal{R}, \mathfrak{W})$ and $(\mathcal{S}, \mathfrak{W})$ are $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{S}_p \mathcal{S}$, they are non-empty and by theorem 3.3, they are mutually exclusive. Then the $\mathcal{S}_t \mathcal{N} \alpha^* g - d$ isconnection of $(\tau_{\mathcal{R}} \mathcal{X}, \underline{\mathbb{U}}, \mathfrak{W})$ presents a contradiction. Hence $(\underline{\mathbb{U}}, \mathfrak{W})$ is not a combination of any two $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{S}_p \mathcal{S}$.

Conversely, assume that $(\underline{\mathbb{U}},\underline{\mathfrak{W}})$ is not a combination of any two $\mathcal{S}_t\mathcal{N}\alpha^*g-\mathcal{S}_ps$. Suppose $(\tau_{\mathcal{R}}\mathcal{X},\underline{\mathbb{U}},\underline{\mathfrak{W}})$ is $\mathcal{S}_t\mathcal{N}\alpha^*g-\mathcal{D}\mathcal{C}_n$. Then $(\underline{\mathbb{U}},\underline{\mathfrak{W}})=(\mathcal{R},\underline{\mathfrak{W}})\cup(\mathcal{S},\underline{\mathfrak{W}})$ where $(\mathcal{R},\underline{\mathfrak{W}})$ and $(\mathcal{S},\underline{\mathfrak{W}})$ are two mutually exclusive non-empty $\mathcal{S}_t\mathcal{N}\alpha^*g-\mathcal{O}s$. Since $(\mathcal{R},\underline{\mathfrak{W}})=(\underline{\mathbb{U}},\underline{\mathfrak{W}})\setminus(\mathcal{S},\underline{\mathfrak{W}})$ and $(\mathcal{S},\underline{\mathfrak{W}})=(\underline{\mathbb{U}},\underline{\mathfrak{W}})\setminus(\mathcal{R},\underline{\mathfrak{W}})$, $(\mathcal{R},\underline{\mathfrak{W}}),(\mathcal{S},\underline{\mathfrak{W}})\in\mathcal{S}_t\mathcal{N}\alpha^*g\mathcal{C}(\tau_{\mathcal{R}}\mathcal{X},\underline{\mathbb{U}},\underline{\mathfrak{W}})$. Then $(\mathcal{R},\underline{\mathfrak{W}})\cap\mathcal{S}_t\mathcal{N}\alpha^*g\mathcal{C}(\mathcal{S},\underline{\mathfrak{W}})=(\mathcal{R},\underline{\mathfrak{W}})\cap\mathcal{S}_t\mathcal{N}\alpha^*g\mathcal{C}((\underline{\mathbb{U}},\underline{\mathfrak{W}})\setminus(\mathcal{R},\underline{\mathfrak{W}}))=(\mathcal{R},\underline{\mathfrak{W}})\cap((\underline{\mathbb{U}},\underline{\mathfrak{W}})\setminus(\mathcal{R},\underline{\mathfrak{W}}))=\emptyset$. Similarly, $\mathcal{S}_t\mathcal{N}\alpha^*g\mathcal{C}(\mathcal{R},\underline{\mathfrak{W}})\cap(\mathcal{S},\underline{\mathfrak{W}})=\emptyset$. Thus $(\mathcal{R},\underline{\mathfrak{W}})$ and $(\mathcal{S},\underline{\mathfrak{W}})$ are $\mathcal{S}_t\mathcal{N}\alpha^*g-\mathcal{S}_p s$, which leads to a contradiction of our assumption. Hence $(\tau_{\mathcal{R}}\mathcal{X},\underline{\mathbb{U}},\underline{\mathfrak{W}})$ is a $\mathcal{S}_t\mathcal{N}\alpha^*g-\mathcal{C}_n\mathcal{S}$.

Theorem 4.10: If $(\mathcal{R}, \mathfrak{W})$ is $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{C}_n$ and $(\mathcal{S}, \mathfrak{W}), (\mathcal{T}, \mathfrak{W})$ are $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{S}_p s$ such that $(\mathcal{R}, \mathfrak{W}) \subset_{\mathcal{S}_t \mathcal{N}} (\mathcal{S}, \mathfrak{W}) \cup (\mathcal{T}, \mathfrak{W})$, then $(\mathcal{R}, \mathfrak{W})$ lies entirely in $(\mathcal{S}, \mathfrak{W})$ or in $(\mathcal{T}, \mathfrak{W})$.

Proof: Suppose $(\mathcal{R},\mathfrak{W}) \not\subset_{\mathcal{S}_t\mathcal{N}} (\mathcal{S},\mathfrak{W})$ and $(\mathcal{R},\mathfrak{W}) \not\subset_{\mathcal{S}_t\mathcal{N}} (\mathcal{T},\mathfrak{W})$. Let $(\mathcal{R}_1,\mathfrak{W}) = (\mathcal{R},\mathfrak{W}) \cap (\mathcal{S},\mathfrak{W})$ and $(\mathcal{R}_2,\mathfrak{W}) = (\mathcal{R},\mathfrak{W}) \cap (\mathcal{T},\mathfrak{W})$. Since $(\mathcal{R}_1,\mathfrak{W}) \subseteq_{\mathcal{S}_t\mathcal{N}} (\mathcal{S},\mathfrak{W})$, $(\mathcal{R}_2,\mathfrak{W}) \subseteq_{\mathcal{S}_t\mathcal{N}} (\mathcal{T},\mathfrak{W})$ and $(\mathcal{S},\mathfrak{W}),(\mathcal{T},\mathfrak{W})$ are $\mathcal{S}_t\mathcal{N}\alpha^*g - \mathcal{S}_p\mathcal{S}$, $(\mathcal{R}_1,\mathfrak{W}) \cap \mathcal{S}_t\mathcal{N}\alpha^*g cl(\mathcal{R}_2,\mathfrak{W}) \subseteq_{\mathcal{S}_t\mathcal{N}} (\mathcal{S},\mathfrak{W}) \cap \mathcal{S}_t\mathcal{N}\alpha^*g cl(\mathcal{T},\mathfrak{W}) = \emptyset$. Similarly, $\mathcal{S}_t\mathcal{N}\alpha^*g cl(\mathcal{R}_1,\mathfrak{W}) \cap (\mathcal{R}_2,\mathfrak{W}) = \emptyset$. Thus $(\mathcal{R}_1,\mathfrak{W})$ and $(\mathcal{R}_2,\mathfrak{W})$ are $\mathcal{S}_t\mathcal{N}\alpha^*g - \mathcal{S}_p\mathcal{S}$. Since $(\mathcal{R},\mathfrak{W}) \subset_{\mathcal{S}_t\mathcal{N}} (\mathcal{S},\mathfrak{W}) \cup (\mathcal{T},\mathfrak{W})$, $(\mathcal{R}_1,\mathfrak{W}) \cup (\mathcal{R}_2,\mathfrak{W}) = [(\mathcal{R},\mathfrak{W}) \cap (\mathcal{S},\mathfrak{W})] \cup [(\mathcal{R},\mathfrak{W}) \cap (\mathcal{S},\mathfrak{W})] \cup (\mathcal{R}_2,\mathfrak{W})$ where $(\mathcal{R}_1,\mathfrak{W})$ and $(\mathcal{R}_2,\mathfrak{W})$ are $\mathcal{S}_t\mathcal{N}\alpha^*g - \mathcal{S}_p\mathcal{S}$. Then by theorem 4.9, $(\mathcal{R},\mathfrak{W})$ is $\mathcal{S}_t\mathcal{N}\alpha^*g - \mathcal{D}\mathcal{C}_n$, leading to a contradiction. Thus either $(\mathcal{R},\mathfrak{W}) \subset_{\mathcal{S}_t\mathcal{N}} (\mathcal{S},\mathfrak{W})$ or $(\mathcal{R},\mathfrak{W}) \subset_{\mathcal{S}_t\mathcal{N}} (\mathcal{S},\mathfrak{W})$.

Theorem 4.11: If $(\mathcal{R}, \mathfrak{W})$ is $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{C}_n$ and $(\mathcal{R}, \mathfrak{W}) \subset_{\mathcal{S}_t \mathcal{N}} (\mathcal{S}, \mathfrak{W}) \subset_{\mathcal{S}_t \mathcal{N}} (\mathcal{S}, \mathfrak{W}) \subset_{\mathcal{S}_t \mathcal{N}} (\mathcal{S}, \mathfrak{W})$, then $(\mathcal{S}, \mathfrak{W})$ is $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{C}_n$.

Proof: Suppose $(\mathcal{S}, \mathfrak{B})$ is $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{D} \mathcal{C}_n$. Then by theorem 4.9, there exists two $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{S}_{\mathcal{P}} \mathcal{S}(\mathcal{S}_1, \mathfrak{B}), (\mathcal{S}_2, \mathfrak{B})$ such that $(\mathcal{S}, \mathfrak{B}) = (\mathcal{S}_1, \mathfrak{B}) \cup (\mathcal{S}_2, \mathfrak{B})$. Since $(\mathcal{R}, \mathfrak{B}) \subset_{\mathcal{S}_t \mathcal{N}} (\mathcal{S}, \mathfrak{B}) = (\mathcal{S}_1, \mathfrak{B}) \cup (\mathcal{S}_2, \mathfrak{B})$, by theorem 4.10, $(\mathcal{R}, \mathfrak{B}) \subset_{\mathcal{S}_t \mathcal{N}} (\mathcal{S}_1, \mathfrak{B})$ or $(\mathcal{R}, \mathfrak{B}) \subset_{\mathcal{S}_t \mathcal{N}} (\mathcal{S}_2, \mathfrak{B})$. we can assume, without limiting generality that $(\mathcal{R}, \mathfrak{B}) \subset_{\mathcal{S}_t \mathcal{N}} (\mathcal{S}_1, \mathfrak{B})$. Then $\mathcal{S}_t \mathcal{N} \alpha^* gcl(\mathcal{R}, \mathfrak{B}) \cap (\mathcal{S}_2, \mathfrak{B}) \subset_{\mathcal{S}_t \mathcal{N}} \mathcal{S}_t \mathcal{N} \alpha^* gcl(\mathcal{S}_1, \mathfrak{B}) \cap (\mathcal{S}_2, \mathfrak{B}) = \emptyset$. Since $(\mathcal{S}_1, \mathfrak{B}) \cup (\mathcal{S}_2, \mathfrak{B}) = (\mathcal{S}, \mathfrak{B}) \subset_{\mathcal{S}_t \mathcal{N}} \mathcal{S}_t \mathcal{N} \alpha^* gcl(\mathcal{R}, \mathfrak{B}), (\mathcal{S}_2, \mathfrak{B}) \cap \mathcal{S}_t \mathcal{N} \alpha^* gcl(\mathcal{R}, \mathfrak{B}) = (\mathcal{S}_2, \mathfrak{B})$. Thus $(\mathcal{S}_2, \mathfrak{B}) = \emptyset$ which is a contradiction, since $(\mathcal{S}_2, \mathfrak{B})$ is non-empty. Similarly, by taking $(\mathcal{R}, \mathfrak{B}) \subset_{\mathcal{S}_t \mathcal{N}} (\mathcal{S}_2, \mathfrak{B})$, we encounter a contradiction to $(\mathcal{S}_1, \mathfrak{B}) \neq \emptyset$. Hence $(\mathcal{S}, \mathfrak{B})$ is $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{C}_n$.

Theorem 4.12: If $(\mathcal{R}, \mathfrak{W})$ is $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{C}_n$, then $\mathcal{S}_t \mathcal{N} \alpha^* g c l(\mathcal{R}, \mathfrak{W})$ is $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{C}_n$.

Proof: Suppose $\mathcal{S}_t \mathcal{N} \alpha^* gcl(\mathcal{R}, \mathfrak{W})$ is $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{D} \mathcal{C}_n$. Then by theorem 4.9, $\mathcal{S}_t \mathcal{N} \alpha^* gcl(\mathcal{R}, \mathfrak{W}) = (\mathcal{R}_1, \mathfrak{W}) \cup (\mathcal{R}_2, \mathfrak{W})$ where $(\mathcal{R}_1, \mathfrak{W})$ and $(\mathcal{R}_2, \mathfrak{W})$ are $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{S}_p \mathcal{S}$. Since $(\mathcal{R}, \mathfrak{W}) \subset_{\mathcal{S}_t \mathcal{N}} \mathcal{S}_t \mathcal{N} \alpha^* gcl(\mathcal{R}, \mathfrak{W}) = (\mathcal{R}_1, \mathfrak{W}) \cup (\mathcal{R}_2, \mathfrak{W})$, by theorem 4.10, either $(\mathcal{R}, \mathfrak{W}) \subset_{\mathcal{S}_t \mathcal{N}} (\mathcal{R}_1, \mathfrak{W})$ or $(\mathcal{R}, \mathfrak{W}) \subset_{\mathcal{S}_t \mathcal{N}} (\mathcal{R}_2, \mathfrak{W})$. If $(\mathcal{R}, \mathfrak{W}) \subset_{\mathcal{S}_t \mathcal{N}} (\mathcal{R}_1, \mathfrak{W})$, then $(\mathcal{R}_2, \mathfrak{W}) \subset_{\mathcal{S}_t \mathcal{N}} (\mathcal{R}_1, \mathfrak{W}) \cup (\mathcal{R}_2, \mathfrak{W}) = \mathcal{S}_t \mathcal{N} \alpha^* gcl(\mathcal{R}, \mathfrak{W}) \subset_{\mathcal{S}_t \mathcal{N}} \mathcal{S}_t \mathcal{N} \alpha^* gcl(\mathcal{R}_1, \mathfrak{W})$. Also since $(\mathcal{R}_1, \mathfrak{W})$ and $(\mathcal{R}_2, \mathfrak{W})$ are $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{S}_p \mathcal{S}_t (\mathcal{R}_1, \mathfrak{W}) \cap (\mathcal{S}_t \mathcal{N} \alpha^* gcl(\mathcal{R}_1, \mathfrak{W})) \cap (\mathcal{R}_2, \mathfrak{W}) = \emptyset$. Then $(\mathcal{R}_2, \mathfrak{W}) \subseteq_{\mathcal{S}_t \mathcal{N}} (\mathcal{V}, \mathfrak{W}) \setminus_{\mathcal{S}_t \mathcal{N}} \mathcal{N} \alpha^* gcl(\mathcal{R}_1, \mathfrak{W})$. Thus $(\mathcal{R}_2, \mathfrak{W}) \subset_{\mathcal{S}_t \mathcal{N}} \mathcal{S}_t \mathcal{N} \alpha^* gcl(\mathcal{R}_1, \mathfrak{W}) \cap ((\mathcal{V}, \mathfrak{W}) \setminus_{\mathcal{S}_t \mathcal{N}} \alpha^* gcl(\mathcal{R}_1, \mathfrak{W})) = \emptyset$ which leas to a contradiction, since $(\mathcal{R}_2, \mathfrak{W})$ is nonempty. Similarly, if we consider $(\mathcal{R}, \mathfrak{W}) \subseteq_{\mathcal{S}_t \mathcal{N}} (\mathcal{R}_2, \mathfrak{W})$, then we encounter a contradiction to $(\mathcal{R}_1, \mathfrak{W}) \neq \emptyset$. Thus $\mathcal{S}_t \mathcal{N} \alpha^* gcl(\mathcal{R}, \mathfrak{W})$ is $\mathcal{S}_t \mathcal{N} \alpha^* gcl(\mathcal{R}, \mathfrak{W}) \subseteq_{\mathcal{S}_t \mathcal{N}} (\mathcal{R}_2, \mathfrak{W})$, then we encounter a contradiction to $(\mathcal{R}_1, \mathfrak{W}) \neq \emptyset$. Thus $\mathcal{S}_t \mathcal{N} \alpha^* gcl(\mathcal{R}, \mathfrak{W})$ is $\mathcal{S}_t \mathcal{N} \alpha^* gcl(\mathcal{R}, \mathfrak{W}) \subseteq_{\mathcal{S}_t \mathcal{N}} (\mathcal{R}_2, \mathfrak{W})$, then we encounter a contradiction to $(\mathcal{R}_1, \mathfrak{W}) \neq \emptyset$.

Remark 4.13: If $(\mathcal{R}, \mathfrak{W}) \subset_{\mathcal{S}_t \mathcal{N}} (\mathcal{S}, \mathfrak{W}) \subset_{\mathcal{S}_t \mathcal{N}} (\mathcal{T}, \mathfrak{W})$ such that $(\mathcal{R}, \mathfrak{W})$ and $(\mathcal{T}, \mathfrak{W})$ are both $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{C}_n$, then $(\mathcal{S}, \mathfrak{W})$ is not necessarily $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{C}_n$.

Example 4.14: Let $U = \{i, j, k, \ell\}$, $\mathfrak{B} = \{m_1, m_2, m_3\}$ and $X = \{i, j\} \subseteq U$ with $U \setminus \mathcal{R} = \{\{i, j\}, \{k, \ell\}\}$. Then $S_t \mathcal{N} \alpha^* g \mathcal{O}(\tau_{\mathcal{R}} X, U, \mathfrak{B}) = \{\emptyset, (U, \mathfrak{B}), (K_2, \mathfrak{B}), (K_3, \mathfrak{B}), (K_6, \mathfrak{B}), (K_{12}, \mathfrak{B}), (K_{13}, \mathfrak{B})\}$. Even though (K_2, \mathfrak{B}) and (K_1, \mathfrak{B}) are $S_t \mathcal{N} \alpha^* g - C_n$ and $(K_2, \mathfrak{B}) \subset_{S_t \mathcal{N}} (K_6, \mathfrak{B}) \subset_{S_t \mathcal{N}} (K_1, \mathfrak{B})$, (K_1, \mathfrak{B}) is $S_t \mathcal{N} \alpha^* g - \mathcal{D} C_n$.





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Theorem 4.15: The combination of any two $S_t \mathcal{N} \alpha^* g - \mathcal{C}_n s$ that have non-empty intersection is $S_t \mathcal{N} \alpha^* g - \mathcal{C}_n$. **Proof:** Let($\mathcal{R}, \mathfrak{W}$) and (S, \mathfrak{W}) be any two $S_t \mathcal{N} \alpha^* g - \mathcal{C}_n s$ such that ($\mathcal{R}, \mathfrak{W}$) \cap (S, \mathfrak{W}) \neq \emptyset . Suppose ($\mathcal{R}, \mathfrak{W}$) \cup (S, \mathfrak{W}) is $S_t \mathcal{N} \alpha^* g - \mathcal{D} \mathcal{C}_n$. Then by theorem 4.9, there exists two $S_t \mathcal{N} \alpha^* g - S_p s(\mathcal{R}_1, \mathfrak{W})$, (S_1, \mathfrak{W}) such that ($\mathcal{R}, \mathfrak{W}$) \cup (S, \mathfrak{W}) = ($\mathcal{R}_1, \mathfrak{W}$) \cup (S_1, \mathfrak{W}). Since ($\mathcal{R}_1, \mathfrak{W}$) and (S_1, \mathfrak{W}) are $S_t \mathcal{N} \alpha^* g - S_p s$, (S_1, \mathfrak{W}) and (S_1, \mathfrak{W}) are non-empty and by theorem 3.3, (S_1, \mathfrak{W}) \cap (S_1, \mathfrak{W}) = \emptyset . Since (S_1, \mathfrak{W}) \cap (S_1, \mathfrak{W}) \cap (S_1, \mathfrak{W}) = \emptyset . Since (S_1, \mathfrak{W}) \cap (S_1, \mathfrak{W}) \cap (S_1, \mathfrak{W}) \cap (S_1, \mathfrak{W}). Also since (S_1, \mathfrak{W}) \cap (S_1, \mathfrak{W}) \cap (S_1, \mathfrak{W}) \cap (S_1, \mathfrak{W}). Also since (S_1, \mathfrak{W}) \cap (S_1, \mathfrak{W}) \cap (S_1, \mathfrak{W}) and (S_1, \mathfrak{W}) and (S_1, \mathfrak{W}) \cap (S_1, \mathfrak{W}) and (S_1, \mathfrak{W}) \cap (S_1, \mathfrak{W}) and (S_1, \mathfrak{W}) \cap (S_1, \mathfrak{W}) and (S_1, \mathfrak{W}) \cap (S_1, \mathfrak{W}) and (S_1, \mathfrak{W}) \cap (S_1, \mathfrak{W}) is non-empty.

Case ii) If $(\mathcal{R}, \mathfrak{W}) \subset_{\mathcal{S}_t \mathcal{N}} (\mathcal{S}_1, \mathfrak{W})$ and $(\mathcal{S}, \mathfrak{W}) \subset_{\mathcal{S}_t \mathcal{N}} (\mathcal{S}_1, \mathfrak{W})$, then $(\mathcal{R}, \mathfrak{W}) \cup (\mathcal{S}, \mathfrak{W}) = (\mathcal{S}_1, \mathfrak{W})$ and $(\mathcal{R}_1, \mathfrak{W}) = \emptyset$ which is a contradiction to $(\mathcal{R}_1, \mathfrak{W})$ is non-empty.

Case iii) If $(\mathcal{R}, \mathfrak{W}) \subset_{\mathcal{S}_{t} \mathcal{N}} (\mathcal{R}_{1}, \mathfrak{W})$ and $(\mathcal{S}, \mathfrak{W}) \subset_{\mathcal{S}_{t} \mathcal{N}} (\mathcal{S}_{1}, \mathfrak{W})$, then $(\mathcal{R}, \mathfrak{W}) \cap (\mathcal{S}, \mathfrak{W}) \subset_{\mathcal{S}_{t} \mathcal{N}} (\mathcal{R}_{1}, \mathfrak{W}) \cap (\mathcal{S}_{1}, \mathfrak{W}) = \emptyset$ which leads to a contradiction of our assumption.

Case iv) If $(\mathcal{R}, \mathfrak{W}) \subset_{\mathcal{S}_t \mathcal{N}} (\mathcal{S}_1, \mathfrak{W})$ and $(\mathcal{S}, \mathfrak{W}) \subset_{\mathcal{S}_t \mathcal{N}} (\mathcal{R}_1, \mathfrak{W})$, then $(\mathcal{R}, \mathfrak{W}) \cap (\mathcal{S}, \mathfrak{W}) \subset_{\mathcal{S}_t \mathcal{N}} (\mathcal{S}_1, \mathfrak{W}) \cap (\mathcal{R}_1, \mathfrak{W}) = \emptyset$ which leads to a contradiction of our assumption. Hence $(\mathcal{R}, \mathfrak{W}) \cup (\mathcal{S}, \mathfrak{W})$ is $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{C}_n$.

Soft nano α^* -generalized hyperconnectedness

Hyper connected spaces lack disconnected components, playing a crucial role in understanding strong connectivity properties and the structure of certain topological spaces. This segment introduces a novel form of hyper connectedness in soft nano topology namely soft nano α^* -generalized hyperconnectedness, elucidating its distinctive properties and implications for soft nano topological space analysis.

Definition 5.1: A subset $(\mathcal{R}, \mathfrak{W})$ of a $\mathcal{S}_t \mathcal{N}TS(\tau_{\mathcal{R}} \mathcal{X}, \mathcal{V}, \mathfrak{W})$ is said to be

- 1) $S_t \mathcal{N} \alpha^* g$ -dense if $S_t \mathcal{N} \alpha^* gcl(\mathcal{R}, \mathfrak{W}) = (U, \mathfrak{W})$
- 2) $S_t \mathcal{N} \alpha^* g$ -nowhere dense if $S_t \mathcal{N} \alpha^* gint(S_t \mathcal{N} \alpha^* gcl(\mathcal{R}, \mathfrak{B})) = \emptyset$

Definition 5.2: A $\mathcal{S}_t \mathcal{N} TS(\tau_{\mathcal{R}} \mathcal{X}, \mathbf{U}, \mathfrak{B})$ is called soft nano α^* -generalized hyperconnected $(\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{H} \mathcal{C}_n)$ if every non-empty $\mathcal{S}_t \mathcal{N} \alpha^* g - 0s$ of $(\tau_{\mathcal{R}} \mathcal{X}, \mathbf{U}, \mathfrak{B})$ is $\mathcal{S}_t \mathcal{N} \alpha^* g$ —dense.

Example 5.3: Let $U = \{i, j, k, \ell\}$, $\mathfrak{W} = \{m_1, m_2, m_3\}$ and $X = \{i, k\} \subseteq U$ with $U \setminus \mathbb{R} = \{\{k\}, \{j, \ell\}\}$. Then $S_t \mathcal{N} \alpha^* g \mathcal{O}(\tau_{\mathcal{R}} X, U, \mathfrak{W}) = \{\emptyset, (U, \mathfrak{W}), (K_1, \mathfrak{W}), (K_2, \mathfrak{W}), (K_1, \mathfrak{W}), (K_{11}, \mathfrak{W}), (K_{12}, \mathfrak{W}), (K_{14}, \mathfrak{W}), (K_{15}, \mathfrak{W})\}$. Since every non-empty $S_t \mathcal{N} \alpha^* g - 0s$ of $(\tau_{\mathcal{R}} X, U, \mathfrak{W})$ is $S_t \mathcal{N} \alpha^* g$ -dense, $(\tau_{\mathcal{R}} X, U, \mathfrak{W})$ is $S_t \mathcal{N} \alpha^* g - \mathcal{H} \mathcal{C}_n$.

Theorem 5.4: For any $S_t \mathcal{N}TS(\tau_R \mathcal{X}, U, \mathfrak{W})$, the subsequent assertions are equivalent.

- i) $(\tau_{\mathcal{R}} \mathcal{X}, \mathcal{Y}, \mathfrak{Y})$ is $\mathcal{S}_t \mathcal{N} \alpha^* g \mathcal{H} \mathcal{C}_n$.
- i) For every soft nano set $(\mathcal{R}, \mathfrak{W})$ of $(\tau_{\mathcal{R}} \mathcal{X}, \mathcal{Y}, \mathfrak{W})$, $(\mathcal{R}, \mathfrak{W})$ is either $\mathcal{S}_t \mathcal{N} \alpha^* g$ —dense or $\mathcal{S}_t \mathcal{N} \alpha^* g$ —nowhere dense.
- ii) Any pair of non-empty $S_t \mathcal{N} \alpha^* g O \operatorname{sof} (\tau_{\mathcal{R}} \mathcal{X}, \mathbf{U}, \mathfrak{B})$ has non-empty intersection.

Proof: i) \Rightarrow ii) Let($\mathcal{R}, \mathfrak{W}$) be a soft nano set of $(\tau_{\mathcal{R}} \mathcal{X}, \cup{U}, \mathfrak{W})$, which is not $\mathcal{S}_t \mathcal{N} \alpha^* g$ -nowhere dense. Then $\mathcal{S}_t \mathcal{N} \alpha^* gint(\mathcal{S}_t \mathcal{N} \alpha^* gcl(\mathcal{R}, \mathfrak{W})) \neq \emptyset$. Since $(\tau_{\mathcal{R}} \mathcal{X}, \cup{U}, \cup{W})$ is $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{H} \mathcal{C}_n$ and $\mathcal{S}_t \mathcal{N} \alpha^* gint(\mathcal{S}_t \mathcal{N} \alpha^* gcl(\mathcal{R}, \cup{W})) \in \mathcal{S}_t \mathcal{N} \alpha^* gol(\tau_{\mathcal{R}} \mathcal{X}, \cup{U}, \cup{W})$, is $\mathcal{S}_t \mathcal{N} \alpha^* g - \text{dense}$. Then $\mathcal{S}_t \mathcal{N} \alpha^* gcl(\mathcal{S}_t \mathcal{N} \alpha^* gcl(\mathcal{S}_t \mathcal{N} \alpha^* gcl(\mathcal{R}, \cup{W})) = (\cup{U}, \cup{W}) \subseteq \mathcal{S}_t \mathcal{N} \alpha^* gcl(\mathcal{R}, \cup{W})$, which implies $\mathcal{S}_t \mathcal{N} \alpha^* gcl(\mathcal{R}, \cup{W}) = (\cup{U}, \cup{W})$

 $S_t \mathcal{N} \alpha^* gcl(S_t \mathcal{N} \alpha^* gint(S_t \mathcal{N} \alpha^* gcl(\mathcal{R}, \mathfrak{W}))) = (U, \mathfrak{W}) \subseteq_{S_t \mathcal{N}} S_t \mathcal{N} \alpha^* gcl(\mathcal{R}, \mathfrak{W}), \text{ which implies } S_t \mathcal{N} \alpha^* gcl(\mathcal{R}, \mathfrak{W}) = (U, \mathfrak{W})$ and hence $(\mathcal{R}, \mathfrak{W})$ is $S_t \mathcal{N} \alpha^* g$ -dense.

ii) \Rightarrow iii) Suppose($\mathcal{R}, \mathfrak{W}$) and ($\mathcal{S}, \mathfrak{W}$) be any two non-empty $\mathcal{S}_t \mathcal{N} \alpha^* g - Os$ of $(\tau_{\mathcal{R}} \mathcal{X}, \mathbf{U}, \mathfrak{W})$ such that ($\mathcal{R}, \mathfrak{W}$) \cap ($\mathcal{S}, \mathfrak{W}$) = \emptyset . Then $\mathcal{S}_t \mathcal{N} \alpha^* gcl(\mathcal{R}, \mathfrak{W}) \cap (\mathcal{S}, \mathfrak{W}) = \emptyset$. Since ($\mathcal{S}, \mathfrak{W}$) \neq \emptyset , $\mathcal{S}_t \mathcal{N} \alpha^* gcl(\mathcal{R}, \mathfrak{W}) \neq$ ($\mathcal{U}, \mathfrak{W}$) and hence ($\mathcal{R}, \mathfrak{W}$) is not $\mathcal{S}_t \mathcal{N} \alpha^* g - dense$. Since ($\mathcal{R}, \mathfrak{W}$) \in $\mathcal{S}_t \mathcal{N} \alpha^* gol(\tau_{\mathcal{R}} \mathcal{X}, \mathcal{U}, \mathfrak{W})$,

 $(\mathcal{R}, \mathfrak{W}) = \mathcal{S}_t \mathcal{N} \alpha^* gint(\mathcal{R}, \mathfrak{W}) \subseteq_{\mathcal{S}_t \mathcal{N}} \mathcal{S}_t \mathcal{N} \alpha^* gint(\mathcal{S}_t \mathcal{N} \alpha^* gcl(\mathcal{R}, \mathfrak{W})).$ Since $(\mathcal{R}, \mathfrak{W}) \neq \emptyset$, $\mathcal{S}_t \mathcal{N} \alpha^* gint(\mathcal{S}_t \mathcal{N} \alpha^* gcl(\mathcal{R}, \mathfrak{W})) \neq \emptyset$. Then $(\mathcal{R}, \mathfrak{W})$ is not $\mathcal{S}_t \mathcal{N} \alpha^* g$ —nowhere dense. Hence $(\mathcal{R}, \mathfrak{W})$ is neither $\mathcal{S}_t \mathcal{N} \alpha^* g$ — dense nor $\mathcal{S}_t \mathcal{N} \alpha^* g$ —nowhere dense, which encounter contradiction to our assumption.





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iii) \Rightarrow i) Suppose($\tau_{\mathcal{R}}\mathcal{X}, \underline{\mathbb{U}}, \mathfrak{B}$) is not $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{H} \mathcal{C}_n$. Then there exists a non-empty $\mathcal{S}_t \mathcal{N} \alpha^* g - Os(\mathcal{T}, \mathfrak{B})$ of $(\tau_{\mathcal{R}}\mathcal{X}, \underline{\mathbb{U}}, \mathfrak{B})$, which is not $\mathcal{S}_t \mathcal{N} \alpha^* g - d$ ense. Then $\mathcal{S}_t \mathcal{N} \alpha^* g cl(\mathcal{T}, \mathfrak{B}) \neq (\underline{\mathbb{U}}, \mathfrak{B})$. Thus $(\underline{\mathbb{U}}, \mathfrak{B}) \setminus \mathcal{S}_t \mathcal{N} \alpha^* g cl(\mathcal{T}, \mathfrak{B})$ and $(\mathcal{T}, \mathfrak{B})$ are non-empty $\mathcal{S}_t \mathcal{N} \alpha^* g - Os$ of $(\tau_{\mathcal{R}}\mathcal{X}, \underline{\mathbb{U}}, \mathfrak{B})$ such that $((\underline{\mathbb{U}}, \mathfrak{B}) \setminus \mathcal{S}_t \mathcal{N} \alpha^* g cl(\mathcal{T}, \mathfrak{B})) \cap (\mathcal{T}, \mathfrak{B}) = \emptyset$, which encounter contradiction to our assumption. Hence $(\tau_{\mathcal{R}}\mathcal{X}, \underline{\mathbb{U}}, \mathfrak{B})$ is $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{H} \mathcal{C}_n$.

Remark 5.5: In a $S_t \mathcal{N}TS(\tau_{\mathcal{R}}\mathcal{X}, \underline{U}, \mathfrak{W})$, if $S_t \mathcal{N}\alpha^* gO(\tau_{\mathcal{R}}\mathcal{X}, \underline{U}, \mathfrak{W}) = \{\emptyset, (\underline{U}, \mathfrak{W})\}$, then $(\tau_{\mathcal{R}}\mathcal{X}, \underline{U}, \mathfrak{W})$ is $S_t \mathcal{N}\alpha^* g - \mathcal{H}C_n$.

Theorem 5.6: If $\mathcal{S}_t \mathcal{N} \alpha^* g \mathcal{O}(\tau_{\mathcal{R}} \mathcal{X}, \c{U}, \cents{m}) = \{ \emptyset, (\c{U}, \c{M}), (\c{R}, \c{M}) \}$, then $(\tau_{\mathcal{R}} \mathcal{X}, \c{U}, \c{M})$ is $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{H} \mathcal{C}_n$. **Proof:** Since (\c{U}, \c{M}) and (\c{R}, \c{M}) are the only non-empty $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{O}s$ of $(\tau_{\mathcal{R}} \mathcal{X}, \c{U}, \c{M}), (\c{U}, \c{M}) \cap (\c{R}, \c{M}) = (\c{R}, \c{M}) \neq \emptyset$. Then by theorem 5.4, $(\tau_{\mathcal{R}} \mathcal{X}, \c{U}, \c{M})$ is $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{H} \mathcal{C}_n$.

Theorem 5.7: Every $S_t \mathcal{N} \alpha^* g - \mathcal{H} C_n S$ is $S_t \mathcal{N} \alpha^* g - C_n$.

Proof: Let $(\tau_{\mathcal{R}}\mathcal{X}, \c{U}, \centsymbol{\mathfrak{B}})$ be $\mathcal{S}_t\mathcal{N}\alpha^*g - \mathcal{H}\mathcal{C}_n$. Then by theorem 5.4, the intersection of any two non-empty $\mathcal{S}_t\mathcal{N}\alpha^*g - \mathcal{S}_t\mathcal{N}\alpha^*g - \mathcal{S}_t\mathcal{$

Remark 5.8: The reverse of the aforementioned theorem may not hold true, as demonstrated by the subsequent example.

Example 5.9: Let $U = \{i, j, k, \ell\}$, $\mathfrak{W} = \{m_1, m_2, m_3\}$ and $\mathcal{X} = \{j, \ell\} \subseteq U$ with $U \setminus \mathcal{R} = \{\{i\}, \{j\}, \{k, \ell\}\}$. Then $S_t \mathcal{N} \alpha^* g \mathcal{O}(\tau_{\mathcal{R}} \mathcal{X}, U, \mathfrak{W}) = \{\emptyset, (U, \mathfrak{W}), (K_3, \mathfrak{W}), (K_4, \mathfrak{W}), (K_5, \mathfrak{W}), (K_9, \mathfrak{W}), (K_{10}, \mathfrak{W}), (K_{11}, \mathfrak{W}), (K_{15}, \mathfrak{W})\}$. Here $(\tau_{\mathcal{R}} \mathcal{X}, U, \mathfrak{W})$ is $S_t \mathcal{N} \alpha^* g - C_n$ but not $S_t \mathcal{N} \alpha^* g - \mathcal{H} C_n$, since only (U, \mathfrak{W}) and (K_{15}, \mathfrak{W}) are $S_t \mathcal{N} \alpha^* g$ -dense.

Theorem 5.10: In a $S_t \mathcal{N}TS(\tau_R \mathcal{X}, \mathcal{U}, \mathfrak{B})$, the subsequent assertions are equivalent.

- a) $(\tau_{\mathcal{R}} \mathcal{X}, \mathcal{V}, \mathfrak{V})$ is $\mathcal{S}_t \mathcal{N} \alpha^* g \mathcal{H} \mathcal{C}_n$.
- b) $S_t \mathcal{N} \alpha^* g$ -interior of every proper $S_t \mathcal{N} \alpha^* g$ Cs of $(\tau_R \mathcal{X}, U, \mathfrak{W})$ is empty.

Proof: a) \Rightarrow b) Let $(\tau_{\mathcal{R}}\mathcal{X}, \cupsyset{\mathbb{U}}, \cupsyset{\mathbb{W}})$ is $S_t \mathcal{N} \alpha^* g - \mathcal{H} \mathcal{C}_n$. If $(\mathcal{R}, \cupsyset{\mathbb{W}})$ is a proper $S_t \mathcal{N} \alpha^* g - Os$ in $(\tau_{\mathcal{R}}\mathcal{X}, \cupsyset{\mathbb{U}}, \cupsyset{\mathbb{W}})$, then $(\mathcal{R}, \cupsyset{\mathbb{W}})^c$ is a proper $S_t \mathcal{N} \alpha^* g - Cs$ in $(\tau_{\mathcal{R}}\mathcal{X}, \cupsyset{\mathbb{U}}, \cupsyset{\mathbb{W}})$. By remark 2.7 a), $S_t \mathcal{N} \alpha^* gint(\mathcal{R}, \cupsyset{\mathbb{W}})^c = (S_t \mathcal{N} \alpha^* gcl(\mathcal{R}, \cupsyset{\mathbb{W}}))^c$. Since $(\tau_{\mathcal{R}}\mathcal{X}, \cupsyset{\mathbb{U}}, \cupsyset{\mathbb{W}})$ is $S_t \mathcal{N} \alpha^* g - Cs$ in $(\tau_{\mathcal{R}}\mathcal{X}, \cupsyset{\mathbb{U}}, \cupsyset{\mathbb{W}})^c = (\cupsyset{\mathbb{U}}, \cupsyset{\mathbb{W}}) = (\cupsyset{\mathbb{U}}, \cupsyset{\mathbb{W}})$. Therefore, $S_t \mathcal{N} \alpha^* gint(\mathcal{R}, \cupsyset{\mathbb{W}})^c = (\cupsyset{\mathbb{U}}, \cupsyset{\mathbb{W}}) \Rightarrow$ a) Assume that the $S_t \mathcal{N} \alpha^* g$ —interior of every proper $S_t \mathcal{N} \alpha^* g - Cs$ of $(\tau_{\mathcal{R}}\mathcal{X}, \cupsyset{\mathbb{U}}, \cupsyset{\mathbb{W}})$ is a proper $S_t \mathcal{N} \alpha^* g - Cs$ in $(\tau_{\mathcal{R}}\mathcal{X}, \cupsyset{\mathbb{U}}, \cupsyset{\mathbb{W}})$, then $(\cupsyset{\mathcal{R}}, \cupsyset{\mathbb{W}})^c$ is a proper $S_t \mathcal{N} \alpha^* g - Os$ in $(\tau_{\mathcal{R}}\mathcal{X}, \cupsyset{\mathbb{U}}, \cupsyset{\mathbb{W}})^c = (\cupsyset{\mathbb{U}}, \cupsyset{\mathbb{W}})^c$ and by our assumption, $S_t \mathcal{N} \alpha^* gint(\cupsyset{\mathcal{R}}, \cupsyset{\mathbb{W}}) = \cupsyset{\mathbb{U}}$. By remark 2.7 b), $S_t \mathcal{N} \alpha^* gcl(\cupsyset{\mathbb{R}}, \cupsyset{\mathbb{W}})^c = (\cupsyset{\mathbb{U}}, \cupsyset{\mathbb{W}})^c$. Then $S_t \mathcal{N} \alpha^* gcl(\cupsyset{\mathbb{R}}, \cupsyset{\mathbb{W}})^c = (\cupsyset{\mathbb{U}}, \cupsyset{\mathbb{W}})^c$ is a proper $S_t \mathcal{N} \alpha^* gcl(\cupsyset{\mathbb{R}}, \cupsyset{\mathbb{W}})^c = (\cupsyset{\mathbb{U}}, \cupsyset{\mathbb{W}})^c$. Then $S_t \mathcal{N} \alpha^* gcl(\cupsyset{\mathbb{R}}, \cupsyset{\mathbb{W}})^c = (\cupsyset{\mathbb{U}}, \cupsyset{\mathbb{W}})^c$. Then $S_t \mathcal{N} \alpha^* gcl(\cupsyset{\mathbb{R}}, \cupsyset{\mathbb{W}})^c = (\cupsyset{\mathbb{U}}, \cupsyset{\mathbb{W}})^c$. Hence $(\cupsyset{\mathbb{U}}, \cupsyset{\mathbb{U}}, \cupsyset{\mathbb{U}})^c$. Hence $(\cupsyset{\mathbb{U}}, \cupsyset{\mathbb{U}}, \cupsyset{\mathbb{U}})^c$ is a proper $\cupsyset{\mathbb{U}}, \cupsyset{\mathbb{U}}$. Hence $(\cupsyset{\mathbb{U}}, \cupsyset{\mathbb{U}}, \cupsyset{\mathbb{U}})^c$ is a proper $\cupsyset{\mathbb{U}}$

Theorem 5.11: $AS_t \mathcal{N}TS(\tau_{\mathcal{R}} \mathcal{X}, \mathbf{U}, \mathfrak{B})$ is $S_t \mathcal{N} \alpha^* g - \mathcal{H} \mathcal{C}_n$ if and only if the combination of any two non-universal $S_t \mathcal{N} \alpha^* g - \mathcal{C}s$ is not universal.

Proof: Let $(\tau_{\mathcal{R}}\mathcal{X}, \c{U}, \cent{M})$ be $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{H} \mathcal{C}_n$ and $(\mathcal{R}, \cent{M}), (\mathcal{S}, \cent{M})$ be two $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{C} s$ such that $(\mathcal{R}, \cent{M}) \neq (\cup{U}, \cent{M})$ and $(\mathcal{S}, \cent{M}) \neq (\cup{U}, \cent{M})$. Then $(\mathcal{R}, \cent{M})^c$ and $(\mathcal{S}, \cent{M})^c$ are non-empty $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{O} s$. Since $(\tau_{\mathcal{R}} \mathcal{X}, \cup{U}, \cent{M})$ is $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{H} \mathcal{C}_n$, by theorem 5.4, $(\mathcal{R}, \cent{M})^c \cap (\mathcal{S}, \cent{M})^c \neq \emptyset$. This implies $((\mathcal{R}, \cent{M}) \cup (\mathcal{S}, \cent{M}))^c \neq \emptyset$ and hence $(\mathcal{R}, \cent{M}) \cup (\mathcal{S}, \cent{M}) \neq (\cup{U}, \cent{M})$. Conversely, let $(\mathcal{R}, \cent{M}), (\mathcal{S}, \cent{M})$ be two $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{C} s$ such that $(\mathcal{R}, \cent{M}) \neq (\cup{U}, \cent{M})$, $(\mathcal{S}, \cent{M}) \neq (\cup{U}, \cent{M})$ and $(\mathcal{R}, \cent{M}) \cup (\mathcal{S}, \cent{M}) \neq (\cup{U}, \cent{M})$. Then $((\mathcal{R}, \cent{M}) \cup (\mathcal{S}, \cent{M}))^c \neq \emptyset$ which implies $(\mathcal{R}, \cent{M})^c \cap (\mathcal{S}, \cent{M})^c \neq \emptyset$, where $(\mathcal{R}, \cent{M})^c$ and $(\mathcal{S}, \cent{M})^c$ are non-empty $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{H} \mathcal{C}_n$.

Theorem 5.12: $\mathcal{S}_t \mathcal{N} \alpha^* g$ —irresolute surjective image of a $\mathcal{S}_t \mathcal{N} \alpha^* g$ — $\mathcal{HC}_n \mathcal{S}$ is $\mathcal{S}_t \mathcal{N} \alpha^* g$ — \mathcal{HC}_n .

Proof: Let $f: (\tau_{\mathcal{R}} \mathcal{X}, \c{U}, \c{M}) \to (\sigma_{\mathcal{R}} \mathcal{Y}, \c{V}, H)$ be a $\mathcal{S}_t \mathcal{N} \alpha^* g$ –irresolute surjection and $(\tau_{\mathcal{R}} \mathcal{X}, \c{U}, \c{M})$ be $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{H} \mathcal{C}_n$. Suppose $(\sigma_{\mathcal{R}} \mathcal{Y}, \c{V}, H)$ is not $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{H} \mathcal{C}_n$. Then there exists a non-empty $\mathcal{S}_t \mathcal{N} \alpha^* g - Os(\mathcal{R}, H)$ of $(\sigma_{\mathcal{R}} \mathcal{Y}, \c{V}, H)$, which





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is not $\mathcal{S}_t \mathcal{N} \alpha^* g$ -dense, that is $\mathcal{S}_t \mathcal{N} \alpha^* g c l(\mathcal{R}, H) \neq (\mathcal{V}, H)$. Since f is $\mathcal{S}_t \mathcal{N} \alpha^* g$ -irresolute surjection, $f^{-1}(\mathcal{R}, H) \in \mathcal{S}_t \mathcal{N} \alpha^* g O(\tau_{\mathcal{R}} \mathcal{X}, \underline{\mathbb{U}}, \mathfrak{B})$ and by remark 2.9, $\mathcal{S}_t \mathcal{N} \alpha^* g c l(f^{-1}(\mathcal{R}, H)) \subseteq_{\mathcal{S}_t \mathcal{N}} f^{-1}(\mathcal{S}_t \mathcal{N} \alpha^* g c l(\mathcal{R}, H)) \neq f^{-1}(\mathcal{V}, H)$, which implies $\mathcal{S}_t \mathcal{N} \alpha^* g c l(f^{-1}(\mathcal{R}, H)) \neq (\underline{\mathbb{U}}, \mathfrak{B})$. Thus $f^{-1}(\mathcal{R}, H)$ is a non-empty $\mathcal{S}_t \mathcal{N} \alpha^* g - O s$ of $(\tau_{\mathcal{R}} \mathcal{X}, \underline{\mathbb{U}}, \mathfrak{B})$, which is not $\mathcal{S}_t \mathcal{N} \alpha^* g$ -dense. This encounters a contradiction to $(\tau_{\mathcal{R}} \mathcal{X}, \underline{\mathbb{U}}, \mathfrak{B})$ is $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{H} \mathcal{C}_n$. Hence $(\sigma_{\mathcal{R}} \mathcal{Y}, \mathcal{V}, H)$ is $\mathcal{S}_t \mathcal{N} \alpha^* g - \mathcal{H} \mathcal{C}_n$.

CONCLUSION

This article delves into the realm of soft nano topology, shedding light on the significance of soft nano α^* -generalized connected spaces by introducing soft nano α^* -generalized separate sets. Their properties, equivalent conditions, interrelation with relevant examples and the intricacies of their preservation theorem are thoroughly explored. Additionally, the idea of soft nano α^* -generalized hyperconnected spaces are discussed along with its characteristics. Nowadays soft nano sets find diverse applications including machine reasoning, artificial neural networks, and databases for smart computing, among others. Soft nano topology, with its focus on connectedness and hyper connectedness, revolutionizes nanoscience by enabling precise control over molecular interactions, fostering innovations in diverse fields like medicine and electronics.

Conflict of Interest

"The authors declare no conflict of interests".

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RESEARCH ARTICLE

Discrete Labeling of Some Trees

A.Punitha Tharani¹ and P.Saradha^{2*}

¹Associate Professor, Department of Mathematics, St. Mary's, College(Autonomous), Thoothukudi, Tamil Nadu, India, (Affiliated to Manonmaniam Sundaranar University, Abishekapatti, Tirunelveli), Tamil Nadu, India.

²Research Scholar (Register number: 21212212092005), Department of Mathematics, St.Mary's College (Autonomous), Thoothukudi, Tamil Nadu, India, (Affiliated to Manonmaniam Sundaranar University, Abishekapatti, Tirunelveli), Tamil Nadu, India.

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*Address for Correspondence

P.Saradha

Research Scholar (Register number: 21212212092005),

Department of Mathematics, St. Mary's College (Autonomous),

Thoothukudi, Tamil Nadu, India, (Affiliated to Manonmaniam Sundaranar University, Abishekapatti,

Tirunelveli), Tamil Nadu, India. Email: sara1998dha@gmail.com



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ABSTRACT

In this paper we deal with Discrete labeling of some trees. We have previously defined discrete labeling which aims at providing the maximum output whenever the inputs are distinct with rational distribution of neighbours thereby exhibiting the stronger form of cordial labeling. This work serves as an aid in decision making. The discrete labels 0 and 1 are cordially assigned to the vertices such that the edges receive labels depending on the incident vertex labels using EX-OR operation with the condition that for every vertex the cardinality of neighbours labeled 0 and 1 differs by at most 1. This paper proposes the discrete labeling of some trees like H-graph, Perfect binary tree and Banana tree. We discuss various cases under which the above said graphs satisfy discrete labeling.

Keywords: Discrete, neighbours, H-graph, Perfect binary tree, Banana tree.

INTRODUCTION

Labeling of graphs is a function that maps the vertex set (edge set) to the set of labels. Here, the domain and co domain are the set of vertices and {0, 1} respectively. Motivated by the cordial labeling [1], we have previously defined a new type of labeling called "Discrete labeling" [2] which assigns labels using EX-OR operations. In our previous works, we have proved that certain standard graphs like path, star, comb, bistar, complete bipartite, broom and some special graphs are discrete. We have also found that friendship graphs, complete graphs, book graphs are





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not discrete[2][3]. There are also graphs like cycle C_n , wheel $W_{1,n}$ which are discrete only for some n. We have found that certain standard and special graphs are discrete and obtained conditions under which some cycle related graphs admit discrete labeling. In this paper, we have obtained the discrete labeling of some trees. The graph G(V, E)discussed here are simple, connected and undirected. The terminologies and symbols used in this paper are in accordance with [4].

METHODS

Definition 2.1:Let G(V, E) be a simple, connected, undirected graph. G is said to have a discrete labeling if there exist functions $d: V \to \{0,1\}$ and $e: E \to \{0,1\}$ defined by

$$e(uv) = \begin{cases} 0 \ ; d(u) = d(v) \\ 1 \ ; \ d(u) \neq d(v) \end{cases} u, v \in V \text{for which}$$
$$|n_d(0) - n_d(1)| \le 1 \text{ ---(i)}$$

 $|n_e(0) - n_e(1)| \le 1$ and ---(ii)

 $|n_{N(v)}(0) - n_{N(v)}(1)| \le 1 \ \forall \ v \in V, \text{---(iii)}$

where $n_d(x)$ and $n_e(x)$ denote the number of vertices and edges with d(u) = x and e(uv) = x; $x \in \{0,1\}$ respectively and $n_{N(v)}(0)$ and $n_{N(v)}(1)$ denote the number of neighbours of the vertex v labeled 0 and 1 respectively. A Graph G is discrete if it admits discrete labeling.[2]

RESULTS AND DISCUSSION

Theorem 3.1:

H-graph is discrete

Proof: The H-graph is the graph obtained from two copies of path P_n with the vertices $u_1, u_2, ..., u_n$ and $v_1, v_2, ..., v_n$ by joining the vertices $u_{\frac{n+1}{2}}$ and $v_{\frac{n+1}{2}}$ if n is odd and $u_{\frac{n}{2}+1}$ and $v_{\frac{n}{2}}$ if n is even. Thus the H-graph has 2n vertices and 2n-1

Define a vertex function $f: V \to \{0,1\}$ by

$$f(u_1) = 0$$
 $f(u_{2i}) = f(u_{2i+1}) = \begin{cases} 1 & \text{when i is odd} \\ 0 & \text{when i is even} \end{cases}$

Now let us label the vertices v_i 's as follows:

Case 1: When $n \equiv 0 \mod 4$

$$f(v_1) = 0$$

$$f(v_{2i}) = f(v_{2i+1}) = \begin{cases} 1 & \text{when i is odd} \\ 0 & \text{when i is even} \end{cases}$$

Case 2: When $n \equiv 1 \mod 4$

$$f(v_{2i}) = f(v_{2i-1}) = \begin{cases} 1 & \text{when } i \text{ is odd} \\ 0 & \text{when } i \text{ is even} \end{cases}$$

Case 3: When $n \equiv 2 \mod 4$

$$f(v_1) = 1$$

$$f(v_{2i}) = f(v_{2i+1}) = \begin{cases} 0 & \text{when i is odd} \\ 1 & \text{when i is even} \end{cases}$$

Case 4: When $n \equiv 3 \mod 4$

$$f(v_{2i}) = f(v_{2i-1}) = \begin{cases} 0 & \text{when } i \text{ is odd} \\ 1 & \text{when } i \text{ is even} \end{cases}$$

 $f(v_{2i}) = f(v_{2i-1}) = \begin{cases} 0 & \textit{when i is odd} \\ 1 & \textit{when i is even} \end{cases}$ Now label the edges as 1 if the incident vertex labels are distinct and 0 otherwise. The number of vertices and edges labeled 0 and 1 and for every vertex the number of neighbouring vertices labeled 0 and 1 are listed below in Table 1: It is observed from the table that $|n_f(0) - n_f(1)| = 0$, $|n_e(0) - n_e(1)| = 1$, $|n_{N_{v_i}}(0) - n_{N_{v_i}}(1)| \le 1$ for all $1 \le i \le 1$ n and $|n_{N_{u_i}}(0) - n_{N_{u_i}}(1)| \le 1$ for all $1 \le i \le n$. Hence H-graph admits Discrete labeling.





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Theorem 3.2:

Perfect binary tree admits discrete labeling

Proof: A Perfect binary tree is a binary tree in which all the interior nodes have two children and all leaves have the same depth or same level. Hence the number of vertices and edges in a perfect binary tree of height h are $2^{h+1}-1$ and $2^{h+1} - 2$ respectively. Let u denote the root vertex of the Perfect binary tree v_1, v_2 denote the vertices in the first level ; v_{2^1+1} , v_{2^1+2} ..., $v_{2^1+2^2}$ denote the vertices in the second level; $v_{2^1+2^2+1}$, $v_{2^1+2^2+2}$, ..., $v_{2^1+2^2+2^3}$ denote the vertices in the third Proceeding like this, at the level h the vertices level. $v_{2^1+2^2+\cdots+2^{h-1}+1}, v_{2^1+2^2+\cdots+2^{h-1}+2}, \dots, v_{2^1+2^2+\cdots+2^{h-1}+2^h}$. Define a vertex function $f: V \to \{0,1\}$ by

$$f(v_i) = \begin{cases} f(u) = 0 \\ 0 & \text{when } i \text{ is odd} \\ 1 & \text{when } i \text{ is even} \end{cases}$$

and label the edges as 1 if the vertex labels are distinct and 0 otherwise. The number of vertices and edges labeled 0 and 1 and for every vertex the number of neighbouring vertices labeled 0 and 1 are listed below in Table 2:

Hence we shall conclude that $|n_f(0) - n_f(1)| \le 1$, $|n_e(0) - n_e(1)| \le 1$, $|n_{N_{v_i}}(0) - n_{N_{v_i}}(1)| \le 1$ for all i and $|n_{N_u}(0) - n_{N_{v_i}}(0)| \le 1$ $n_{N_{v_{1}}}(1) \mid = 0.$

Theorem 3.3:

Banana Tree $B_{n,k}$ admits discrete labeling only when n=2 for all k and when n=3 and k is even.

Proof: An (n, k) banana tree is a graph obtained by connecting one leaf of each of n copies of a k-star graph with a single root vertex that is distinct from all stars. Thus in $B_{n,k}$, |V| = nk + 1 and |E| = nk. Let w denote the root vertex of $B_{n,k}$, $u_i(1 \le i \le n)$ denote the vertices of the stars attached to the root vertex, $v_i(1 \le i \le n)$ denote the central vertex of the stars and x_{ij} ($1 \le i \le n$ and $1 \le j \le k - 2$) denote the pendant vertices of the n stars respectively.

Necessary part: Case 1: When n = 2 and k is odd($k \ge 3$).

The vertex function $f: V \rightarrow \{0,1\}$ defined by

$$f(w) = 0$$

$$f(u_i) = \begin{cases} 0 \text{ when } i \text{ is odd} \\ 1 \text{ when } i \text{ is even} \end{cases}$$

$$f(v_i) = 1 \ \forall \ i$$

$$f(x_{ij}) = \begin{cases} 0; 1 \le j \le \frac{k-3}{2} \\ 1; \frac{k-3}{2} + 1 \le j \le k - 2 \end{cases}$$
 when *i* is odd
$$f(x_{ij}) = \begin{cases} 1; 1 \le j \le \frac{k-3}{2} \\ 0; \frac{k-3}{2} + 1 \le j \le k - 2 \end{cases}$$
 when *i* is even

satisfies the discrete labeling. Now, assign the label 1 to the edges which are incident with distinct vertex labels and 0 otherwise. The number of vertices, edges and for every vertex the number of neighbours labeled 0 and 1 are tabulated below in Table 3:

Hence it is clear that $|n_f(0) - n_f(1)| = 1$, $|n_e(0) - n_e(1)| = 0$ and for every vertex, the number of neighbours labeled 0 and 1 differs by at most 1 which proves that $B_{n,k}$ when n=2 and k is odd.

Case 2: When n = 2.3 and k is even.

The vertex function $f: V \to \{0,1\}$ defined by

$$f(w) = 0$$

$$f(u_i) = \begin{cases} 0 \text{ when } i \text{ is odd} \\ 1 \text{ when } i \text{ is even} \end{cases}$$

$$f(u_i) = \{1, when i is even$$

$$f(v_i) = 1 \ \forall \ i$$

$$f(x_{ij}) = \begin{cases} 0; 1 \le j \le \frac{k-2}{2} \\ 1; \frac{k-2}{2} + 1 \le j \le k-2 \end{cases}$$

satisfies the discrete labeling. Now, assign the label 1 to the edges which are incident with distinct vertex labels and 0 otherwise. The number of vertices, edges and for every vertex the number of neighbours labeled 0 and 1 are





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tabulated below in Table 4: Hence it is clear that $|n_f(0) - n_f(1)| = 1$, $|n_e(0) - n_e(1)| = 0$ and for every vertex, the number of neighbours labeled 0 and 1 differs by at most 1. Therefore, it can be concluded that $B_{n,k}$ is discrete when n = 2,3 and k is even.

Sufficient Part: Case 3: When k is odd for $n \ge 3$.

W.l.o.g , label the root vertex w as 0 and to maintain the desired absolute difference of vertices labeled 0 and 1 let $f(u_i) = \begin{cases} 0 \text{ when } i \text{ is odd} \\ 1 \text{ when } i \text{ is even} \end{cases}$. Now since u_i is of degree 2 and f(w) = 0, all v_i 's must be assigned the label 1 in order to satisfy $|n_{N(u_i)}(0) - n_{N(u_i)}(1)| \le 1$. At this juncture , we are left with labeling n sets of $k - 2x_{ij}$'s. Keeping in mind that the neighbours of v_i 's are even, we are supposed to give equal number of 0's and 1's to the neighbours of v_i . Now when i is odd ,by the pre assigned vertex label $f(u_i) = 0$, $\frac{k-3}{2}$ vertices of x_{ij} must be labeled 0 and $\frac{k-1}{2}$ vertices of x_{ij} must be labeled 1. Similarly, When i is even, by the pre assigned vertex label $f(u_i) = 1$, $\frac{k-1}{2}$ vertices of x_{ij} must be labeled 0 and $\frac{k-3}{2}$ vertices of x_{ij} must be labeled 1 to satisfy $|n_{N(v_i)}(0) - n_{N(v_i)}(1)| \le 1$. By labeling so, we find that the number of vertices labeled 0 and 1 are $1 + n\left(\frac{k-1}{2}\right)$ and $n + n\left(\frac{k-1}{2}\right)$ making the absolute difference $|n_f(0) - n_f(1)| = |1 - n| \ge |1 - 3| = 2$ violating the definition of discrete labeling. Hence $B_{n,k}$ is not discrete when k is odd for all $n \ge 3$.

Case 4: When k is even for all $n \ge 4$.

First let us label the vertices optimally satisfying $|n_f(0) - n_f(1)| \le 1$ under $f: V \to \{0,1\}$.

Without loss of generality, label w as 0 and $f(u_i) = \begin{cases} 0 \text{ when } i \text{ is odd} \\ 1 \text{ when } i \text{ is even} \end{cases}$. Since u_i 's are of degree two,all v_i 's must be assigned the label 1 to satisfy $|n_{N(u_i)}(0) - n_{N(u_i)}(1)| \le 1$. Now for x_{ij} when i is odd ,label the first $\frac{k-2}{2}$ vertices 0 and next $\frac{k-2}{2}$ vertices 1. When i is even, label the first $\frac{k-4}{2}$ vertices with 1 and the remaining $\frac{k}{2}$ vertices with 0 which is the only possible way to label x_{ij} without violating $|n_f(0) - n_f(1)| \le 1$. We notice that there are $\frac{nk+2}{2}$ vertices are labeled 0 and remaining $\frac{nk}{2}$ vertices are labeled 1 making $|n_f(0) - n_f(1)| = 1$. Now label the edges 1 whenever the vertex labels are distinct and 0 otherwise. On labeling the edges, we find that when n is even , there are $n + \frac{n}{2}(k-3)$ edges labeled 0 and $n + \frac{n}{2}(k-1)$ edges labeled 1 and when n is odd, there are $\frac{n+1}{2}(\frac{k}{2}) + \frac{n-1}{2}(\frac{k-2}{2})$ edges labeled 0 and $\frac{n+1}{2}(\frac{k}{2}) + \frac{n-1}{2}(\frac{k+2}{2})$ edges labeled 1 making the absolute difference n and n-1 respectively which is obviously greater than or equal to 3 since $n \ge 4$ violating the definition. Hence, $B_{n,k}$ is not discrete when k is even for all $n \ge 4$.

CONCLUSION

We have discussed the Discrete labeling of some tree related graphs and found that not all trees are discrete. Further explorations are being done on characterisation of the class of graphs that admit discrete labeling. Also, this work can be extended by applying the adjacency condition of the vertices to the edges resulting in edge discrete labeling[5].

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Table 1: Vertex condition, Edge condition and neighbouring label condition of H- graph

Table 1: Ve	rtex con	iaition,	Table 1: Vertex condition, Edge condition and neighbouring label condition of H- graph							
Cases	$n_f(0)$	$n_f(1)$	$n_e(0)$	$n_e(1)$	$n_{N_{u_1}}(0)$	$n_{N_{u_1}}(1)$	$n_{N_{v_1}}(0)$	$n_{N_{v_1}}(1)$	$n_{N_{u_i}}(0)$ $2 \le i \le n - 1$	$n_{N_{u_i}}(1)$ $2 \le i \le n - 1$
$n = 0 \bmod 8$	n	n	n-1	n	0	1	0	1	1; $i \neq u_{\frac{n}{2}+1}$ 2; $i = u_{\frac{n}{2}+1}$	1; $i \neq u_{\frac{n}{2}+1}$ 1; $i = u_{\frac{n}{2}+1}$
$n = 1 \mod 8$	n	n	n-1	n	0	1	0	1	1; $i \neq u_{\frac{n+1}{2}}$ 1; $i = u_{\frac{n+1}{2}}$	1; $i \neq u_{\frac{n+1}{2}}$ 2; $i = u_{\frac{n+1}{2}}$
$n = 2 \mod 8$	n	n	n-1	n	0	1	1	0	1; $i \neq u_{\frac{n}{2}+1}$ 1; $i = u_{\frac{n}{2}+1}$	1; $i \neq u_{\frac{n}{2}+1}$ 2; $i = u_{\frac{n}{2}+1}$
$n = 3 \mod 8$	n	n	n-1	n	0	1	1	0	1; $i \neq u_{\frac{n+1}{2}}$ 2; $i = u_{\frac{n+1}{2}}$	1; $i \neq u_{\frac{n+1}{2}}$ 1; $i = u_{\frac{n+1}{2}}$
$n = 4 \mod 8$	n	n	n-1	n	0	1	0	1	1; $i \neq u_{\frac{n}{2}+1}$ 1; $i = u_{\frac{n}{2}+1}$	1; $i \neq u_{\frac{n}{2}+1}$ 2; $i = u_{\frac{n}{2}+1}$
$n = 5 \mod 8$	n	n	n-1	n	0	1	0	1	1; $i \neq u_{\frac{n+1}{2}}$ 2; $i = u_{\frac{n+1}{2}}$	1; $i \neq u_{\frac{n+1}{2}}$ 1; $i = u_{\frac{n+1}{2}}$
$n = 6 \mod 8$	n	n	n-1	n	0	1	1	0	1; $i \neq u_{\frac{n}{2}+1}$ 2; $i = u_{\frac{n}{2}+1}$	1; $i \neq u_{\frac{n}{2}+1}$ 1; $i = u_{\frac{n}{2}+1}$
$n = 7 \mod 8$	n	n	n-1	n	0	1	1	0	1; $i \neq u_{\frac{n+1}{2}}$ 2; $i = u_{\frac{n+1}{2}}$	1; $i \neq u_{\frac{n+1}{2}}$ 1; $i = u_{\frac{n+1}{2}}$





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$n_{N_{v_i}}(0)$	$n_{N_{v_i}}(1)$	$n_{N_{u_n}}(0)$	$n_{N_{u_n}}(1)$	$n_{N_{v_n}}(0)$	$n_{N_{v_n}}(1)$
$1; i \neq v_{\frac{n}{2}}$ $2; i = v_{\frac{n}{2}}$	$1; i \neq v_{\frac{n}{2}}$ $1; i = v_{\frac{n}{2}}$	0	1	0	1
1; $i \neq v_{\frac{n+1}{2}}$ 2; $i = v_{\frac{n+1}{2}}$	$1; i \neq v_{\frac{n+1}{2}}$ $1; i = v_{\frac{n+1}{2}}$	1	0	1	0
$1; i \neq v_{\frac{n}{2}}$ $1; i = v_{\frac{n}{2}}$	$1; i \neq v_{\frac{n}{2}}$ $2; i = v_{\frac{n}{2}}$	1	0	0	1
1; $i \neq v_{\frac{n+1}{2}}$ 1; $i = v_{\frac{n+1}{2}}$	$1; i \neq v_{\frac{n+1}{2}}$ $2; i = v_{\frac{n+1}{2}}$	0	1	1	0
$1; i \neq v_{\frac{n}{2}}$ $1; i = v_{\frac{n}{2}}$	$1; i \neq v_{\frac{n}{2}}$ $2; i = v_{\frac{n}{2}}$	0	1	0	1
1; $i \neq v_{\frac{n+1}{2}}$ 1; $i = v_{\frac{n+1}{2}}$	1; $i \neq v_{\frac{n+1}{2}}$ 2; $i = v_{\frac{n+1}{2}}$	1	0	1	0
$1; i \neq v_{\frac{n}{2}}$ $2; i = v_{\frac{n}{2}}$	$1; i \neq v_{\frac{n}{2}}$ $1; i = v_{\frac{n}{2}}$	1	0	0	1
1; $i \neq v_{\frac{n+1}{2}}$ 1; $i = v_{\frac{n+1}{2}}$	1; $i \neq v_{\frac{n+1}{2}}$ 2; $i = v_{\frac{n+1}{2}}$	0	1	1	0

Table 2: Vertex condition, Edge condition and neighbouring label condition of Perfect binary tree

$n_f(0)$	$n_f(1)$	$n_e(0)$	$n_e(1)$	$n_{N_u}(0)$	$n_{N_u}(1)$	$n_{N_{v_1}}(0)$
2^h	$2^{h}-1$	$2^{h}-1$	$2^{h}-1$	1	1	2

$n_{N_{v_1}}(1)$	$n_{N_{v_2}}(0)$	$n_{N_{v_2}}(1)$	$n_{N_{v_{2i}}}(0)$ $2 \le i \le$ $\frac{2^1 + 2^2 + \dots + 2^{h-1}}{2}$	$2 \le i \le \frac{n_{N_{v_{2i}}}(1)}{2^1 + 2^2 + \dots + 2^{h-1}}$
1	2	1	2 ; i even 1, i = 3,5,	1; <i>i even</i> 2; <i>i</i> = 3,5,





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1; i even	0; i even	1; i even	0; i even
0; i odd	1; i odd	0; i odd	1; i odd

Table 3: Cardinality of vertices, edges and neighbours of $B_{n,k}(k \text{ odd}, n = 2)$

				<u> </u>		п,к
Case (k odd)	$n_f(0)$	$n_f(1)$	$n_e(0)$	$n_e(1)$	$n_{N(w)}(0)$	$n_{N(w)}(1)$
n = 2	$\frac{kn}{2}$	$\frac{kn+2}{2}$	$\frac{kn}{2}$	$\frac{kn}{2}$	1	1

Case(k odd)	$n_{N(u_i)}(0)$	$n_{N(u_i)}(1)$	$n_{N(x_{ij})}(0)$	$n_{N(x_{ij})}(1)$
n = 2	1	1	0	1

Case(k odd)	$n_{N(v_i)}(0)$	$n_{N(v_i)}(1)$
n = 2	$\frac{k-1}{}$	$\frac{k-1}{}$
	2	2

Table 4: Cardinality of vertices, edges and neighbours of $B_{n,k}$ (k even, n=2,3)

Case (k even)	$n_f(0)$	$n_f(1)$	$n_e(0)$	$n_e(1)$	$n_{N(w)}(0)$	$n_{N(w)}(1)$
n = 2	$\frac{kn}{2}$	$\frac{kn+2}{2}$	$\frac{kn}{2}$	$\frac{kn}{2}$	1	1
n = 3	$\frac{kn}{2}$	$\frac{kn+2}{2}$	$\frac{kn}{2}$	$\frac{kn}{2}$	2	1

Case(k even)	$n_{N(u_i)}(0)$	$n_{N(u_i)}(1)$	$n_{N(x_{ij})}(0)$	$n_{N(x_{ij})}(1)$
n = 2	1	1	0	1
n = 3	1	1	0	1

Case(k even)	$n_{N(v_i)}(0)$	$n_{N(v_i)}(1)$
n=2 and	k	k-2
i odd	$\overline{2}$	2
n=2 and	k-2	k
i even	2	$\overline{2}$
n = 3 and	k	k-2
i odd	$\overline{2}$	2
n = 3 and	k-2	k
i even	2	$\frac{\overline{2}}{2}$





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RESEARCH ARTICLE

Cardinal Labeling of Some Standard and Special Graphs

A. Punitha Tharani¹ and S. Pratiksha^{2*}

¹Associate Professor, Department of Mathematics, St. Mary's College (Autonomous), Thoothukudi, Tamil Nadu, India, (Affiliated to Manonmaniam Sundaranar University, Abishekapatti, Tirunelveli), Tamil Nadu, India.

²Research Scholar (Register number: 21212212092004), Department of Mathematics, St.Mary's College (Autonomous), Thoothukudi-628 001, Tamil Nadu, India, (Affiliated to Manonmaniam Sundaranar University, Abishekapatti, Tirunelveli), Tamil Nadu, India.

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*Address for Correspondence

S. Pratiksha

Research Scholar (Register number: 21212212092004),

Department of Mathematics, St.Mary's College (Autonomous),

Thoothukudi-628 001, Tamil Nadu, India,

(Affiliated to Manonmaniam Sundaranar University,

Abishekapatti, Tirunelveli), Tamil Nadu, India.

Email: pratikshaprabhu32@gmail.com



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ABSTRACT

Inspired from magic and graceful[1] labelings, we have introduced a new type of labeling wherein we optimally try to label the vertices and edges of a graph using their total count with no repetition under absolute difference operation. We call this as cardinal labeling and refer to those graphs as cardinal. This paper discusses the cardinal labeling for some standard and special graphs.

Keywords: Cardinal, standard graphs, complete graph, complete bipartite graph, special graphs, cardinality.

INTRODUCTION

Graph labeling in general is defined as assignment of elements to the vertices and edges of a graph under a defined function or condition. In our paper, the graphs are simple, connected and undirected and the elements used to label the vertex set (domain) and the edge set are natural numbers. For the graph G with V(G) and E(G) as the set of vertices and edges respectively, we define cardinal labeling and investigate it for certain standard and special graphs. The notations employed here are with reference to [2].





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METHOD

Definition: Let G(V, E) be a simple connected undirected graph. Let |V| = p, |E| = q and let $C = \{1, 2, ..., p + q\}$. G is said to admit cardinal labeling if the one-to-one function f from the vertex set V into the set C generates a one-to-one edge function f^* onto the set C/f(V) defined by $|f(u) - f(v)| = f^*(uv) \ \forall uv \in E$. A graph G is said to be cardinal if it admits cardinal labeling. Figure 1 illustrates the above definition. For this graph, |V| = 4, |E| = 5 and so we have $C = \{1, 2, ..., 9\}$. We could see in the figure that all the members of C are uniquely assigned to the vertices and edges without repetition. Note that the edge labels are the absolute differences of the incident vertex labels. Cardinal labeling takes its origin by partly combination of vertex magic total labeling and graceful labeling. Unlike these two labelings, there is no constant preserved and the vertex and edge labels form a subset of natural numbers; i.e., no vertex or edge receive zero label.

RESULTS AND DISCUSSION

Cardinal Labeling of some Standard graphs

1.1 Theorem: Path P_n satisfies cardinal labeling.

Proof: Let $v_1, v_2, ..., v_n$ be the vertices of the path P_n where |V| = n, |E| = n - 1 and $C = \{1, 2, ..., 2n - 1\}$. Label the vertices v_i (i = 1, 2, ..., n) of P_n by the function $f: V \to C$ given by

$$f(v_i) = \begin{cases} 2n - i & \text{when } i \text{ is odd} \\ i - 1 & \text{when } i \text{ is even} \end{cases}$$

Next, label the edges of P_n by the induced function $f^*: E \to C/f(V)$ given by,

$$f^*(v_i v_{i+1}) = |2n - i - ((i+1) - 1)| = 2(n-i), \quad i = 1, 2, ..., n-1$$

If so done, all the vertices and the edges of P_n obtain distinct |C| = 2n - 1 labels. Hence path P_n is cardinal.

1.2 Theorem: Star $K_{1,m}$ is cardinal.

Proof: Let u denote the central vertex and let v_i (i = 1, 2, ..., m) denote its pendant vertices. Here |V| = m + 1, |E| = m and $C = \{1, 2, ..., 2m + 1\}$. Define $f: V \to C$ by

$$f(u) = 2m + 1$$

$$f(v_i) = 2m + 1 - i, i = 1, 2, ..., m$$

The induced edge function f^* onto the set C/f(V) is given below.

$$f^*(uv_i) = |2m-1-((2m-1)-i)| = i, i = 1,2,...,m-1$$

The resultant edge labels obtained by the absolute difference of the incident vertex labels are all distinct in C. Also vertices u and $v_i (i = 1, ..., m)$ have distinct labels of C that are not given to $\{E\}$. The resulting labeling of the star graph is cardinal. Hence Star $K_{1,m}$ is cardinal.

1.3 Theorem: Cycle graph C_n admits cardinal labeling.

Proof: C_n has n number of vertices and edges. Let u_i (i = 1,2,...,n) be its vertices. We have $C = \{1,2,...,2n\}$. It is possible to cardinally label cycles if we define the vertex function $f: V \to C$ for the odd and even cases of n as follows. Case 1: When n is odd

$$f(u_1) = 2n$$
, $f(u_n) = 2(n-1)$
 $f(u_{2i}) = 2i - 1$, $i = 1, 2, ..., \left[\frac{n}{2}\right]$
 $f(u_{2i-1}) = 2n - (2i - 1)$, $i = 2, 3, ..., \left[\frac{n-1}{2}\right]$

Case 2: When n is even

$$f(u_1) = 2n$$
, $f(u_n) = 2$, $f(u_{2i}) = 2i - 1$, $i = 1, 2, ..., \left[\frac{n-1}{2}\right]$





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$$f(u_{2i-1}) = 2n - (2i-1),$$
 $i = 2,3,..., \left[\frac{n}{2}\right]$

In both the cases, it is possible to get distinct edge labels of C_n if we label the edges using the condition $|f(u_i) - f(u_j)|$, $i \neq j$.

The induced functions $f^*: E \to C/f(V)$ are discussed for the same.

Case (i) When n is odd

$$f^*(u_1u_2) = |2n - (2(1) - 1)| = 2n - 1$$

$$f^*(u_1u_n) = |2n - 2(n - 1)| = 2$$

$$f^*(u_{n-1}u_n) = |(n - 1) - 1 - (2(n - 1))| = n$$

$$f^*(u_iu_{i+1}) = |(i - 1) - (2n - (i + 1))| = |2(i - n)|, \qquad i = 2,3,...,n - 2$$

Case (ii) When n is even

$$f^*(u_1u_2) = |2n - (2 - 1)| = 2n - 1$$

$$f^*(u_1u_n) = |2n - 2| = 2(n - 1)$$

$$f^*(u_{n-1}u_n) = |2n - (n - 1) - 2| = n - 1$$

$$f^*(u_iu_{i+1}) = |i - 1 - (2n - (i + 1))| = 2(n - i), \qquad i = 2, ..., n - 2$$

Thus, all the vertices and edges receive distinct enumerations of the set *C*. Hence it may be concluded that the cycle admits cardinal labeling.

1.4 Theorem: Complete Bipartite graph $K_{m,n}$ satisfies cardinal labeling.

Proof: Let $u_i(i=1,2,...,m)$ and $v_i(i=1,2,...,n)$ be the vertices of $K_{m,n}$. We have |V|=m+n, |E|=mn and $C=\{1,2,...,m+(m+1)n\}$. Let us define the vertex function f to C by

$$f(u_i) = i$$
, $i = 1,2,...,m$
 $f(v_i) = m + (m+1)(n+1-i)$, $i = 1,2,...,n$

The edge labels are generated by simply finding the absolute difference of the respective vertex labels. Down below are the induced edge functions f^* onto C/f(V).

$$f^*(u_iv_j) = |i - (r + (r+1)(s+1-j))| = (r-i) + (r+1)(s+1-j),$$

$$i = 1, 2, ..., r, \qquad j = 1, 2, ..., s$$

On observation, we see that all the vertices and the edges receive definite labels from the set C. From this it becomes obvious that $K_{m,n}$ satisfies cardinal labeling.

1.5 Theorem: Complete graph K_n does not satisfy cardinal labeling for $n \ge 4$.

Proof: Cardinal labeling is true for the case when n < 4. Assume $n \ge 4$.

 K_n has n(=p) vertices and $\frac{n(n-1)}{2}$ (= q) edges, thus $|C| = \frac{n(n+1)}{2}$. Now if we optimally apply the p+q labels to distinct vertices and edges of the graph such that the definition of cardinal labeling is not violated, we see that at least n-3 labels of the set C gets repeated. Since each vertex is adjacent to every other vertex in K_n , in order to obtain distinct edge labels, we have to remove $\Delta-2$ edges from K_n . Otherwise $\Delta-2$ number of labels gets repeated. In either case we arrive at a contradiction. Hence K_n ($n \ge 4$) does not satisfy cardinal labeling.

Cardinal Labeling of some Special graphs[3]

2.1 Bull graph

The bull graph is a cycle C_3 with two disjoint pendant edges which resembles a bull or a ram. It is a unit distance graph with 5 vertices and edges. For the bull graph, $C = \{1, 2, ..., 10\}$. It is possible to label the graph using C as shown in Figure 4.

2.2 Butterfly graph

The butterfly graph is a planar graph obtained by joining two copies of C_3 with a common vertex and is isomorphic to the friendship graph F_2 . It has 5 vertices and 6 edges and so $C = \{1, 2, ..., 11\}$. Distinct vertex and edge labels could be allotted to the graph from C if we label as below (Figure 5)





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2.3 Moser spindle graph

Named after Leo Moser, Moser Spindle graph is a planar unit distance graph with 7 vertices and 11 edges. Here $C = \{1,2,...18\}$. Figure 6 shows the possible way of labeling the graph cardinally.

2.4 Wagner graph

It is a 3-regular non planar graph with 8 vertices and 12 edges. It is the 8-vertex Möbius ladder graph. We have $C = \{1, 2, ..., 20\}$. Figure 7 shows evidently that the graph is cardinal.

2.5 Observation

- i) Petersen graph is not cardinal.
- ii) Herschel graph does not admit cardinal labeling.

General results of Cardinal labeling

3.1 Observation

For a cardinal graph, the maximum label |C| can only be assigned as a vertex label. i.e. it is impossible to obtain label |C| under f^* .

3.2 Result

If all the vertices of a cardinal graph are labelled with odd labels of *C*, then the cardinality of the set *C* is odd.

Proof: By the above observation, maximum label |C| has to be given to one of the vertices of the cardinal graph G. Since all the labels of V are odd, |C| is also odd.

3.3 Theorem

If all the vertices of a cardinal graph G have odd labels of C, then then the cardinality of the set C is odd and G is a tree.

Proof

We prove that |C| is odd and G is a tree when V(G) have odd vertex labels. The first part is evident from Result 3.2. Without loss of generality, we assume that |V(G)| = n, n is odd, so n distinct odd labels of C are required to label every vertex of G odd. Since all the vertices are given odd labels, f^* can assign only distinct even labels of C to the edges of G. Further since G is cardinal and cardinality of the set C is odd, there can only be C 1 distinct even labels of C that could be given to its edges hence |E(G)| = n - 1. Any graph G(n, n - 1) is certainly a tree, hence the result follows. The converse of theorem 3.3 is not true. i.e. i.e., If the cardinality of C is odd and if C is a tree, then all the vertices of C needn't bear odd labels only, for, it is possible to label path C 1 cardinally with a combination of odd and even vertex labels.

Example:

Here P_5 is a tree and |C| = 9 but it could be observed that not all vertices obtain odd labels despite being cardinal (Figure 8).

3.4 Corollary: Any cardinal graph containing a cycle must have at least one even vertex label of *C* in that cycle. **Proof:** We prove that if the cardinal graph has no even vertex labels then the graph cannot have cycles. Take a cardinal graph that has cycles and suppose that all its vertex labels are odd. Then by theorem 3.3, the cardinal graph is a tree contradicting the fact that the graph has cycles.

CONCLUSION

In this work, we have investigated the cardinal labeling for standard graphs like path, star and cycle and some special graphs. A few results have also been obtained as a result of employing the labeling to these graphs. Explorations are being made on some star related graphs and trees. We have planned to further extend our work on graphs obtained by arithmetic and set operations.

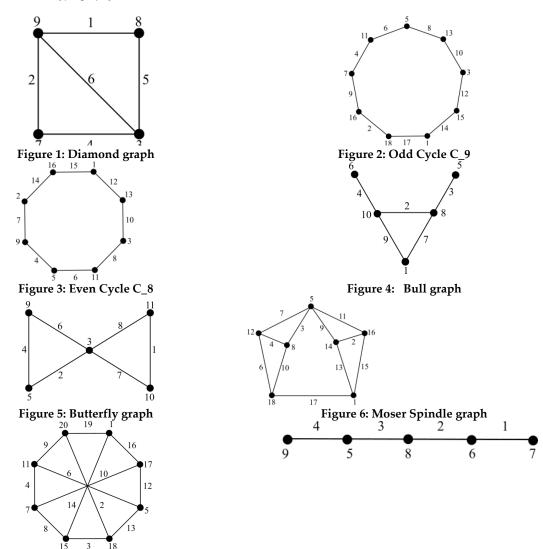




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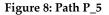




Figure 7: Wagner graph



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RESEARCH ARTICLE

Determinant of Some Cyclic Graphs using Incycle Vertices

Punitha Tharani A1 and Anesha H R2*

¹Associate Professor, Department of Mathematics, St. Mary's College (Autonomous), Thoothukudi, (Affiliated to Manonmaniam Sundaranar University), Abishekapatti, Tirunelveli , Tamil Nadu, India.

²Research Scholar (Reg. No.: 20212212092002), Department of Mathematics, St. Mary's College (Autonomous), Thoothukudi, (Affiliated to Manonmaniam Sundaranar University), Abishekapatti, Tirunelveli, Tamil Nadu, India.

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*Address for Correspondence

Anesha H R

Research Scholar (Reg. No.: 20212212092002),

Department of Mathematics,

St. Mary's College (Autonomous),

Thoothukudi,

(Affiliated to Manonmaniam Sundaranar University),

Abishekapatti, Tirunelveli, Tamil Nadu, India.

Email: aneshahr@gmail.com



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ABSTRACT

The determinant of adjacency matrix of a graph is called the determinant of a graph. Various reduction procedures and formulae have been devised over the years to find the determinant of graphs. Sesquivalent spanning subgraph of a graph is a spanning sub graph in which all components are either cycles or edges. Harary F, gave a formula to find the determinant of a graph using sesquivalent spanning subgraphs. It is an inspiration for the concept of incycle vertices. Perfect matching of a graph, is a matching in which every vertex is covered. Perfect matching is also a sesquivalent spanning subgraph. A vertex is called an incycle vertex if it lies on some cycle in all possible sesquivalent spanning subgraphs of a graph, except the perfect matching. In this paper we have tried to find the determinant of some types of cycle related graphs, using the concept of incycle vertices.

Keywords: Sesquivalent, bicyclic graph, multicyclic graph, determinant, incycle vertex.

INTRODUCTION

Sesquivalent spanning subgraph of a graph is a spanning subgraph in which all components are either cycles or edges. Perfect matching of a graph is a matching in which every vertex is covered. Perfect matching is also a





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sesquivalent spanning subgraph. A vertex is called an *incycle vertex* if it lies on some cycle in all possible sesquivalent spanning subgraphs of a graph, except the perfect matching. [1]If G is a graph whose adjacency matrix is A(G) then $det(A(G)) = \sum (-1)^{r(I)} (2)^{s(I)}$

where $c(\Gamma)$ is the number of components of Γ , $r(\Gamma) = |V(\Gamma)| - c(\Gamma)$ and $s(\Gamma) = |E(\Gamma)| - |V(\Gamma)| + c(\Gamma)$ and the summation is over all sesquivalent spanning subgraphs Γ of G. The above formula was given by Harary F, in 1962. The above formula is an inspiration for the concept of incycle vertices.

METHODS

In this paper we attempt to find the determinant of a few cycle related graphs. ^[2]A unicyclic graph is a tree with an edge that makes a cycle. ^[3]There are 3 types of bicyclic graphs. Consider two vertex disjoint cycles. The graph got by identifying a vertex in one cycle with a vertex of another cycle as v is called a ∞ -graph. Throughout this paper we mean ∞ -graph with trees attached to it when we say bicyclic graph with a common vertex v. We denote the set of all such graphs on v vertices by v vertex from v disjoint cycles. Throughout this paper we mean this graph with trees attached to it when we say multicyclic graph with a common vertex v. We denote the set of all such graphs by v denote the set of all such graphs with only even cycles by v denote the set of all such graphs with only even cycles by v denote the set of all such graphs with only even cycles by v denote the set of all such graphs with only even cycles by v denote the set of all such graphs with only even cycles by v denote the set of all such graphs with only even cycles by v denote the set of all such graphs with only even cycles by v denote the set of all such graphs with only even cycles by v denote the set of all such graphs with only even cycles by v denote the set of all such graphs with only even cycles by v denote the set of all such graphs with only even cycles by v denote the set of all such graphs with only even cycles by v denote the set of all such graphs with only even cycles by v denote the set of all such graphs with only even cycles by v denote the set of all such graphs with only even cycles by v denote the set of all such graphs with only even cycles by v denote the set of all such graphs with even cycles by v denote the set of all such graphs with even cycles by v denote the set of all such graphs with even cycles as v denote the set of all such graphs with even cycles as v denote the set of all such graphs with even cycles as v denote the set of all such gra

Observation 2.1: Incycle vertices always lie on cycles but not all vertices on cycles are incycle vertices. Example: K_n , n > 4. All vertices lie on cycles but no vertex is incycle vertex.

Observation 2.2: All non-singular graphs with cycle need not have an incycle vertex. Example:Sunlet graph. Perfect matching is the only sesquivalent spanning subgraph.

Observation 2.3: Incycle vertices need not be unique.

Example: C_n . All vertices are incycle vertices.

Observation 2.4: Incycle vertices need not lie on every cycle in a graph.

Example: Wheel graph. It is possible to form a cycle with all rim vertices but the central vertex is the only incycle vertex.

Observation 2.5: If v is a vertex that lies on every cycle in a graph, then v belongs to the set of all incycle vertices of the graph.

Observation 2.6: If all cycles have a common vertex, then vertices from two different cycles cannot be incycle, as any sesquivalent spanning subgraph has at most one cycle.

INCYCLE VERTICES IN UNICYCLIC GRAPHS:

Theorem3.1: A unicyclic graph cannot have a unique incycle vertex.

Proof. Suppose a unicyclic graph has a unique incycle vertex, then it lies on a cycle in all sesquivalent spanning subgraphs. But there is only one cycle in a unicyclic graph. Therefore all vertices on that cycle lie on cycles in all sesquivalent spanning subgraphs, which implies all vertices on the cycle are incycle vertices.

Corollary 3.2: If a unicyclic graph has incycle vertices then total number of incycle vertices = the length of the cycle in the unicyclic graph.





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Theorem 3.3: If a unicyclic graph has a sesquivalent spanning subgraph with cycle then all vertices on the cycle are incyclic.

Proof. The unicyclic graph has a sesquivalent spanning subgraph with cycle. The cycle can be odd or even. If the graph contains an odd cycle then no other sesquivalent spanning subgraph is possible. If the graph contains an even cycle then the only other possible sesquivalent spanning subgraphs are perfect matchings. In both cases there are no other cycles in other sesquivalent spanning subgraphs. Therefore all vertices on the cycle are incycle vertices.

Theorem 3.4: If a unicyclic graph has only one sesquivalent spanning subgraph then it has an odd cycle or it is the perfect matching.

Proof. Case 1: The unicyclic graph contains an odd cycle. It can have at most one sesquivalent spanning subgraph. The only possible sesquivalent spanning subgraphs are the perfect matching or one containing the odd cycle in it. Case 2: The unicyclic graph contains an even cycle. The graph has more than one sesquivalent spanning subgraph iff it has a sesquivalent spanning subgraph with the even cycle as one of its components. If there exists a perfect matching in which at least one vertex is matched outside the cycle then no other sesquivalent spanning subgraph is possible.

Note 3.5: The converse is not true for all unicyclic graphs. In case of even unicyclic graph if all vertices in the cycle are matched within itself, in the perfect matching then it has 2 more sesquivalent subgraphs. But if an odd unicyclic graph has a sesquivalent subgraph with odd cycle in it or a perfect matching then it is unique. Also an odd unicyclic graph is non-singular iff it has a sesquivalent spanning subgraph.

Theorem 3.6: A non-singular unicyclic graph has no incycle vertices iff the perfect matching is the only sesquivalent spanning subgraph and |Det(A(G))| = 1.

Proof. Let the unicyclic graph be non-singular. Therefore it has at least one sesquivalent spanning subgraph. Suppose it contains a cycle then by theorem 3.3 all vertices are incycle vertices, which is not possible. Hence the perfect matching is the only sesquivalent spanning subgraph. The converse follows from the definition of incycle vertices.

Theorem 3.7: If a unicyclic graph G has $k \neq 0$ incycle vertices, then

$$Det(A(G)) = \begin{cases} 0, & k \equiv 0 \pmod{4} \\ \pm 4, & k \equiv 2 \pmod{4} \\ \pm 2, & otherwise \end{cases}$$

Proof. Case 1: $k \equiv 0 \pmod{4}$. *G*has 3 sesquivalent spanning subgraphs. One with a cycle and two perfect matchings. $(-1)^{r(\Gamma)}(2)^{s(\Gamma)}$ of the one with the cycle is twice the negative of that of the perfect matchings. Hence their sum is zero. Case 2: $k \equiv 2 \pmod{4}$. *G*has 3 sesquivalent spanning subgraphs. One with a cycle and two perfect matchings. $(-1)^{r(\Gamma)}(2)^{s(\Gamma)}$ of the one with the cycle is twice that of the perfect matchings. Hence their sum is -4 or 4.

Case 3: *k* is odd. G has only one sesquivalent spanning subgraph. It has an odd cycle in it. Hence its determinant is -2 or 2.

In conclusion, if G is a unicyclic graph, $|Det(A(G))| \le 4$.

INCYCLE VERTICES IN A BICYCLIC GRAPHS WITH ONE VERTEX IN COMMON:

Theorem 4.1: If $G \in B_n^{++}$ has an incycle vertex that is not the common vertex then the incycle vertex is not unique. In particular all vertices of that cycle are incyclic.

Proof. If a vertex is an incycle vertex then it lies on a cycle in all sesquivalent spanning subgraphs except the perfect matching. If a vertex other than v is incycle then that cycle is in every sesquivalent spanning subgraph other than the perfect matching. Therefore all its vertices including v are incyclic.

Theorem 4.2: If $G \in B_n^{++}$ has no incyclic vertex then the perfect matching is the only possible sesquivalent spanning subgraph and $|Det(A(G))| \le 1$.





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Proof. If G has a sesquivalent spanning subgraph with cycle in it then at least v is an incycle vertex. If v is not an incycle vertex then G has no incycle vertices. Hence the perfect matching is the only sesquivalent spanning subgraph. If G has a sesquivalent spanning subgraph then its determinant is either -1 or 1. If G has no sesquivalent spanning subgraphs then its determinant is 0.

Corollary 4.3: If $G \in B_n^{++}$ has no incycle vertices then determinant is 0,1 or -1

Theorem 4.4: Let $G \in B_n^{++}$ and let G have no incyclic vertices. Then G is singular iff it has no sesquivalent spanning subgraph.

Proof. Let G be singular. From theorem 4.2G has at most one sesquivalent spanning subgraph. If G has a perfect matching then G is not singular. The converse is obvious.

Corollary 4.5: Let $G \in B_n^{++}$ and let G have no incyclic vertices. Then G is singular if it has odd number of vertices.

Theorem 4.6:If $G \in B_n^{++}$, has $k(\neq 0)$ incycle vertices, then

$$Det(A(G)) = \begin{cases} 0, & k \equiv 0 \pmod{4} \\ \pm 4, & k \equiv 2 \pmod{4} \text{ or } k = 1 \\ \pm 2, & otherwise \end{cases}$$

Proof. Case 1: $k \equiv 0 \pmod{4}$. Ghas 3 sesquivalent spanning subgraphs. One with an even cycle and two perfect matchings. $(-1)^{r(\Gamma)}(2)^{s(\Gamma)}$ of the one with the cycle is twice the negative of that of the perfect matchings. Hence their sum is zero.

Case 2: $k \equiv 2 \pmod{4}$. Ghas 3 sesquivalent spanning subgraphs. One with an even cycle and two perfect matchings. $(-1)^{r(I)}(2)^{s(I)}$ of the one with the cycle is twice that of the perfect matchings. Hence their sum is -4 or 4.

Case 3: $k \neq 1$ is odd. Ghas only one sesquivalent spanning subgraph. It has an odd cycle in it. Hence its determinant is -2 or 2.

Case 4: k = 1. Ghas 2 sesquivalent spanning subgraphs, both with an odd cycle in it. $(-1)^{r(\Gamma)}(2)^{s(\Gamma)}$ for both are the same and their sum is -4 or 4.

Corollary 4.7: If $G \in B_n^{++}$ has an odd number of incycle vertices, then G is non-singular. In conclusion, if $G \in B_n^{++}$, $|Det(A(G))| \le 4$.

INCYCLE VERTICES IN A MULTICYCLIC GRAPHS WITH ONE VERTEX IN COMMON:

Theorem 5.1: If $G \in M_n^{++}$ has no incycle vertex then the perfect matching is the only possible sesquivalent spanning subgraph and $|Det(A(G))| \le 1$.

Proof. If G has a sesquivalent spanning subgraph with cycle in it then at least v is an incycle vertex. If v is not an incycle vertex then G has no incycle vertices. Hence the perfect matching is the only sesquivalent spanning subgraph. If G has a sesquivalent spanning subgraph then its determinant is either -1 or 1. If G has no sesquivalent spanning subgraphs then its determinant is G.

Corollary 5.2: If $G \in M_n^{++}$ has no incycle vertices then determinant is 0,1 or -1

Theorem 5.3: Let $G \in M_n^{++}$ and let G - v have no perfect matching. Then G has sesquivalent spanning subgraph iff there exists a maximum matching of G - v in which exactly one neighbor of v is unmatched.

Proof. Let G have a sesquivalent spanning subgraph. If it is a perfect matching then exactly one vertex was unmatched in G - v and it must be a neighbour of v. If the sesquivalent spanning subgraph has a cycle, the cycle could not be odd. If the cycle is even then all vertices of that cycle should have been matched within itself except one. It is possible to get a maximum matching of G - v in which exactly one neighbor of v is unmatched. Conversely, let there exist a maximum matching of G - v in which exactly one neighbor of v is unmatched. Then by matching it with v in G we get a perfect matching that is a sesquivalent spanning subgraph.

Corollary 5.4: $G \in M_n^{++}$ is singular if in the maximum matching of G - v there are more than one unmatched vertices. **Proof.** By theorem 5.3, G has no sesquivalent spanning subgraph.





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Theorem 5.5: Let $G \in M_{n,}^{++}$. If G - v has no perfect matching and G has odd number of vertices then G is singular. **Proof.** If G has odd number of vertices, G - v has even number of vertices. As G - v has no perfect matching at least 2 vertices are unmatched. By corollary 5.4, G is singular.

Multicyclic graphs having a vertex in common with only Even cycles:

Theorem 5.1.1: Let $G \in M_{n,e}^{++}$. If G - v has a perfect matching then G has no sesquivalent spanning subgraphs and is singular.

Proof. If G - v has a perfect matching then in every cycle of G at least one vertex is matched outside the cycle. Therefore G has no sesquivalent spanning subgraphs. Hence G is singular.

Theorem5.1.2: If $G \in M_{n,e}^{++}$ has incycle vertices. Then G - v has no perfect matching.

Proof. Suppose G - v has a perfect matching. Then G has no sesquivalent spanning subgraph and hence no incycle vertices, which is not possible.

Theorem 5.1.3: If $G \in M_{n,e}^{++}$ has incycle vertex then G has 3 sesquivalent spanning subgraphs.

Proof. *G* has incycle vertex. Then *G* has sesquivalent spanning subgraph with cycle. Consider a sesquivalent spanning subgraph with cycle. By removing the alternating edges in the cycle we get 2 perfect matchings. No other sesquivalent spanning subgraph with cycle is possible as all cycles are even.

Theorem 5.1.4: $G \in M_{n,e}^{++}$ cannot have exactly one incycle vertex. Number of incycle vertices is at least the number of vertices in the smallest cycle in G.

Proof. Suppose G has an incycle vertex. Then by theorem 5.1.3, G has exactly 3sesquivalent spanning subgraphs. Only one of it has a cycle in it. Therefore all the vertices in the cycle are incycle vertices. Hence G has either no incycle vertex or more than one incycle vertices.

Theorem 5.1.5: If $G \in M_{n,e}^{++}$ has $k(\neq 0)$ incycle vertices, then

$$Det(A(G)) = \begin{cases} 0, & k \equiv 0 \pmod{4} \\ \pm 4, & k \equiv 2 \pmod{4} \end{cases}$$

Proof. Case 1: $k \equiv 0 \pmod{4}$. By theorem 5.1.3, G has exactly 3 sesquivalent spanning subgraphs, one with cycle and 2 perfect matchings. One with an even cycle and two perfect matchings. $(-1)^{r(\Gamma)}(2)^{s(\Gamma)}$ of the one with the cycle is twice the negative of that of the perfect matchings. Hence their sum is zero.

Case 2: $k \equiv 2 \pmod{4}$. By theorem 5.1.3, G has exactly 3 sesquivalent spanning subgraphs. One with an even cycle and two perfect matchings. $(-1)^{r(\Gamma)}(2)^{s(\Gamma)}$ for the one with the cycle is twice that of the perfect matchings. Hence their sum is -4 or 4.

Theorem 5.1.6: Let $G \in M_{n,e}^{++}$. If G is non-singular, having incycle vertex then |det(A(G))| = 4.

Proof. G cannot have odd number of incycle vertices as all cycles are even, by theorem 5.1.4. By theorem 5.1.5, if the number of vertices in the cycle is a multiple of 4 then the determinant is 0. But G is non-singular, hence the determinant is -4 or 4.

In conclusion if $G \in M_{n,e}^{++}$, $|Det(A(G))| \le 4$.

Multicyclic graphs having a vertex in common with only Odd cycles:

Theorem 5.2.1: If $G \in M_{n,o}^{++}$ has a perfect matching then G has exactly one sesquivalent spanning subgraph and |det(A(G))| = 1.

Proof. *G* cannot have another perfect matching as all cycles in G are odd. G cannot have sesquivalent spanning subgraphs with cycles in it, as there is no cycle which is matched within itself in the perfect matching. Therefore G has exactly one perfect matching and its determinant is -1 or 1.

Theorem 5.2.2: Let $G \in M_{n,o}^{++}$. Then either all sesquivalent spanning subgraphs of G have cycle in it or G has a perfect matching.





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Proof. From previous theorem if *G* has a perfect matching, it has no other sesquivalent spanning subgraphs. If *G* has a sesquivalent spanning subgraph with cycle in it, then *G* cannot have a perfect matching as all cycles are odd.

Corollary 5.2.3: Let $G \in M_n^{++}$. If G has sesquivalent spanning subgraph with odd cycle in it then, G has no perfect matching.

Theorem 5.2.4: If $G \in M_{n,o}^{++}$ has more than one incycle vertices then G has exactly one sesquivalent spanning subgraph. **Proof.** G has more than one incycle vertices. All vertices of at most one cycle are incycle. All cycles are odd, so if there exists a sesquivalent spanning subgraph with cycle then there is no perfect matching. Also there is at most one sesquivalent spanning subgraph corresponding to a cycle. So G has exactly one sesquivalent spanning subgraph.

Corollary 5.2.5: $G \in M_{n,o}^{++}$ has more than one sesquivalent spanning subgraphs iff G has exactly one incycle vertex. **Proof.** Let G have more than one sesquivalent spanning subgraphs. Then all sesquivalent spanning subgraphs have cycle in it, by theorem 5.2.2. But no two sesquivalent spanning subgraphs have the same cycle in it and hence G has only one incycle vertex. Conversely, let G have exactly one incycle vertex. Then G has at least two sesquivalent spanning subgraphs with cycle in it, corresponding to different cycles.

Theorem 5.2.6: If $G \in M_{n,o}^{++}$ has more than one incycle vertices then G is non-singular and |Det(A(G))| = 2.. **Proof.** By theorem 5.2.4, G has exactly one sesquivalent spanning subgraph. $(-1)^{r(\Gamma)}(2)^{s(\Gamma)}$ for it is -2 or 2. Hence G is non-singular.

Theorem5.2.7: Let $G \in M_{n,o}^{++}$. G is non-singulariff it has at least one sesquivalent spanning subgraph.

Proof. Let G be non-singular. Then G has at least one sesquivalent spanning subgraph. Conversely, let G have a sesquivalent spanning subgraph. If G has a perfect matching G cannot have any other sesquivalent spanning subgraphs. Then G is non-singular. If G has sesquivalent spanning subgraphs with cycle, then $(-1)^{r(f)}(2)^{s(f)}$ is of the same sign for all such subgraphs and hence their sum is greater than zero.

Theorem 5.2.8: Let $G \in M_{n,o}^{++}$ and let G - v have a perfect matching. G is non-singulariff it has at least one incycle vertex.

Proof. Let G be non-singular. As G - v has a perfect matching, G cannot have a perfect matching. By theorem 5.2.7, G has at least one sesquivalent spanning subgraph with cycle. Hence G has at least one incycle vertex. Conversely, let G have at least one incycle vertex. Then G has sesquivalent spanning subgraph with cycle. By theorem 5.2.7, G is non-singular.

Corollary 5.2.9: $G \in M_{n,o}^{++}$ is non-singular if G has incycle vertex.

Theorem 5.2.10: Let $G \in M_{n,o}^{++}$ and let G - v have a perfect matching. *G* is non-singulariff there exists at least one cycle in which all vertices are matched within itself in G - v.

Proof. Let G be non-singular. As G-v has a perfect matching, G cannot have a perfect matching. By theorem 5.2.7, G has at least one sesquivalent spanning subgraph with cycle, which is possible only if at least one cycle is matched within itself. Conversely, let there exist at least one cycle in which all vertices are mapped within itself. Then G has at least one incycle vertex. By corollary 5.2.9, G is non-singular.

Theorem 5.2.11: Let $G \in M_{n,o}^{++}$. If G is non-singular and G - v has a perfect matching then G has exactly x sesquivalent spanning subgraphs, where x is the number of cycles that are matched within themselves in G - v.

Proof. G is non-singular, so G has sesquivalent spanning subgraph by theorem 5.2.7. G-v has a perfect matching, so all sesquivalent spanning subgraphs have cycle in them. If a cycle is matched within itself in G-v then G has a sesquivalent spanning subgraph with that cycle in it. In the same way G has a sesquivalent spanning subgraph with every cycle matched within itself in G-v. Hence the number of sesquivalent spanning subgraphs in G is equal to the number of cycles matched within itself in G-v.





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Corollary 5.2.11:Let $G \in M_{n,o}^{++}$ and let G - v have a perfect matching. If G is non-singular then det(A(G)) is $\pm 2(x)$, where x is the number of cycles that are matched within themselves in G - v.

Proof. From previous theorem, *G* has *x* sesquivalent spanning subgraphs and all of them have cycle in it. $(-1)^{r(I)}(2)^{s(I)}$ is either all -2 or 2 for every such subgraph. Their sum is $\pm 2(x)$.

Theorem 5.2.12: Let $G \in M_{n,o}^{++}|det(A(G))| > 2iff G$ has exactly one incycle vertex.

Proof. Let |det(A(G))| > 2. Suppose G has more than one incycle vertices, then by theorem 5.2.6, |det(A(G))| = 2. Conversely, let G have exactly one incycle vertex. Then by corollary 5.2.5, G has more than one sesquivalent spanning subgraphs, all with cycles in it. Each of the subgraph contributes -2 or 2 to the determinant. They are all of the same sign and hence |det(A(G))| > 2.

Multicyclic graphs having a vertex in common with both Odd and Even cycles:

Theorem 5.3.1: If $:G \in M_n^{++}$ has even number of incycle vertices then G has exactly 3 sesquivalent spanning subgraphs.

Proof. *G* has an even number of incycle vertices. Then *G* has a sesquivalent spanning subgraph with an even cycle. Consider that subgraph. By removing the alternating edges in that cycle we get 2 perfect matchings. No sesquivalent spanning subgraph with any other cycle is possible.

Theorem 5.3.2: Let $G \in M_n^{++}$ has $k \neq 0$ incycle vertices, then

$$Det(A(G)) = \begin{cases} 0, & k \equiv 0 \pmod{4} \\ \pm 4, & k \equiv 2 \pmod{4} \end{cases}$$

Proof. Case 1: $k \equiv 0 \pmod{4}$. By theorem 5.3.1, *G* has exactly 3 sesquivalent spanning subgraphs, one with cycle and 2 perfect matchings. One with an even cycle and two perfect matchings. $(-1)^{r(\Gamma)}(2)^{s(\Gamma)}$ for the one with the cycle is twice the negative of that of the perfect matchings. Hence their sum is zero.

Case 2: $k \equiv 2 \pmod{4}$. By theorem 5.3.1, G has exactly 3 sesquivalent spanning subgraphs. One with an even cycle and two perfect matchings. $(-1)^{r(\Gamma)}(2)^{s(\Gamma)}$ for the one with the cycle is twice that of the perfect matchings. Hence their sum is -4 or 4.

Corollary 5.3.3: If a non-singular $G \in M_n^{++}$ has even number of incycle vertices then $|\det(A(G))| = 4$.

Theorem 5.3.4: If $G \in M_n^{++}$ has an odd number of incycle vertices greater than 1 then |det(A(G))| = 2.

Proof. Ghas a sesquivalent spanning subgraph with an odd cycle and no other sesquivalent spanning subgraph with cycles in it. Gdoes not have a perfect matching. Hence the determinant is -2 or 2.

Theorem 5.3.5: If $G \in M_n^{++}$ has only one incycle vertex then G has more than one odd cycles.

Proof. Ghasonly one incycle vertex. Then G has at least two sesquivalent spanning subgraphs, all with distinct cycles in it. If a sesquivalent spanning subgraph has an even cycle in it, then it is the only sesquivalent spanning subgraph with cycle in it, by theorem 5.3.1. Therefore G has at least 2 odd cycles.

Theorem 5.3.6: Let $G \in M_n^{++}$. If |det(A(G))| > 4, then G has more than two odd cycles.

Proof. From theorem 5.3.2 and theorem 5.3.4 it is evident that G has exactly one incycle vertex. G does not have any sesquivalent spanning subgraph with even cycle in it. G has no perfect matching either. By theorem 5.3.5, G has more than one odd cycles. Suppose G has 2 odd cycles, then G has at most 2 sesquivalent spanning subgraphs, all with distinct cycles in it. Each of the subgraph contributes -2 or 2 to the determinant. They are all of the same sign and hence $|det(A(G))| \le 4$, which is not possible. Hence G has more than two odd cycles.

CONCLUSION

In this paper we have introduced the concept of incycle vertices. We have calculated the determinant of some cyclic graphs using incycle vertices. We arrive at the conclusion that $|det(A(G))| \le 4$, where G is a multicyclic graph with



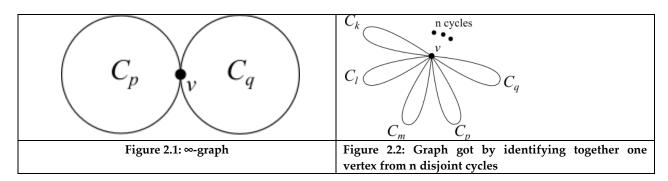


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one vertex in common, having less than 3 odd cycles. Future study will be focused on solving the determinant of all types of bicyclic graphs and other graphs of similar structure.

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RESEARCH ARTICLE

Some New Families of Sum Divisor Cordial Graphs

K. Periasamy^{1*} K. Venugopal¹ and P. Lawrence Rozario Raj²

¹P.G. and Research Department of Mathematics, Dr. Kalaignar Government Arts College, Kulithalai, (Affiliated to Bharathidasan University), Tiruchirappalli, Tamil Nadu, India.

²P.G. and Research Department of Mathematics, St. Joesph's College (Affiliated to Bharathidasan University), Tiruchirappalli, Tamil Nadu, India.

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*Address for Correspondence

K. Periasamy

P.G. and Research Department of Mathematics,

Dr. Kalaignar Government Arts College,

Kulithalai,

(Affiliated to Bharathidasan University),

Tiruchirappalli, Tamil Nadu, India.

Email: kpsperiasamy87@gmail.com.



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ABSTRACT

The eccentric graph of any graph G is a graph constructed from the vertices of G such that any two vertices are joined by an edge only when any one of them is an eccentric vertex of the other. It is denoted by G_{ecc} . In this paper, we investigate the sdc-labeling of $(K_n(e)K_n)_{ecc}$, $(S(K_{1,n})(e)S(K_{1,n})_{ecc}$, $(S(K_{1,n})(e)K_m)_{ecc}$ and $(C_n(e)C_n)_{ecc}$.

Keywords: Eccentric graph, sum divisor cordial graph, complete graph, cycle, subdivision of star graph.

INTRODUCTION

We start with an undirected, simple, finite graph. A graph G = (V, E), where V is the set of vertices with |V| = p and Eis the set of edges with |E| = q, is referred to as a (p,q) graph G. The origins of graph theory are attributed to Leonhard Euler's 1736 examination of the well-known Königsberg bridge problem. Graph labeling is one of the main topics of graph theory. Graph labeling is the process of giving vertices, edges, or both of a graph labels, which are often represented by integers. One of the fascinating and active subfields of graph theory is labeling of graphs. Rosa presented the idea of grace in [10]. Graham et al. presented the idea of harmony in [6]. The graph labeling techniques in graph theory originated with graceful labeling and harmonic labeling. Another significant labeling style that incorporates elements of both graceful and harmonious is called cordial, and it was first established by Cahit in [2]. The idea of divisor cordial labeling of graphs was first presented by Varatharajan et al. in [12]. Lourdusamy et al. presented the idea of sum divisor cordial labeling in [8]. Eccentric graphs are one type of eccentricity-related graph study; these studies of graphs are fascinating and amazing. Akiyama et al. defined the unique kind of graph





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operation known as the eccentric graph of graph in [1], and they also provided several intriguing characterizations of these kinds of graphs. There is only one graph structure driven by any graph action. However, nearly every graph has an eccentric graph operation that leads to multiple fascinating and amazing graph structures. Kaspar et al. discovered and described such intriguing graph topologies of eccentric graphs of specific classes of graphs in [7]. Ghosh et al. in[9] look into a few families of prime cordial eccentric graphs. The graphs under discussion are undirected, simple, and finite. The terms used in graph theory are referred to by Harary [5]. We refer to Tom M. Apostol [11] for the conventional terminology and notations in number theory, and Gallian [3] for the literature on graph labeling. Investigating the sdc-labeling of $(K_n(e)K_n)_{ecc}$, $(S(K_{1,n})(e)S(K_{1,n})_{ecc}$, $(S(K_{1,n})(e)K_m)_{ecc}$, and $(C_n(e)C_n)_{ecc}$ is the goal of this work.

Preliminaries

We will give brief summary of definitions which are useful for the present investigations.

Definition: A graph labeling is the assignment of unique identifiers to the edges and vertices of a graph.

Definition : Let G be a graph with vertex set V and edge set E. The eccentric graph of G is denoted by $G_{ecc} = (V, E')$, where $E' = \{(u,v) : d_G(u,v) = ecc(u) \text{ or } d_G(u,v) = ecc(v) \text{ for } u,v \in V\}$.

Definition : Let G = (V,E) be a simple graph and $f : V(G) \to \{1, 2, ..., |V(G)|\}$ be a bijection. For each edge uv, assign the label 1 if $2 \mid (f(u)+f(v))$ and the label 0 otherwise. The function f is called a sum divisor cordial labeling if $|e_f(0) - e_f(1)| \le 1$. A graph which admits a sum divisor cordial labeling (sdc-labeling) is called a sum divisor cordial graph (sdc-graph).

Definition: An edge joining of two copies of any graph G is denoted by G(e)G.

Definition: An edge joining of two different graph G and H is denoted by G(e)H.

Main Results

Theorem 3.1 $(K_n(e)K_n)_{ecc}$ is sdc-graph for $n \ge 3$.

Proof.

Let G be $K_n(e)K_n$. G has 2n vertices and n^2-n+1 edges. Then the vertices of G are $v_1,v_2,v_3,...,v_n$, $w_1,w_2,w_3,...,w_n$. Let $(K_n(e)K_n)_{ecc}$ be the eccentric graph of $K_n(e)K_n$.

Then $(K_n(e)K_n)_{ecc}$ is $K_1 + \overline{K_{n-1}} + \overline{K_{n-1}} + K_1$, the vertices are v_1 , w_2 , w_3 , ..., w_n , v_2 , v_3 , ..., v_n , v_n , v_n . In $(K_n(e)K_n)_{ecc}$, v_1 is adjacent to v_2 , v_3 , ..., v_n , and also w_2 , w_3 , ..., v_n and v_2 , v_3 , ..., v_n are mutually adjacent. Then, $|V((K_n(e)K_n)_{ecc})| = 2n$ and $|E((K_n(e)K_n)_{ecc})| = n^2 - 1$.

Define $f: V((K_n(e)K_n)_{ecc}) \rightarrow \{1, 2, ..., 2n\}$ as follows.

$$f(v_1) = 2$$
 and $f(w_1) = 1$,

$$f(w_i) = i + n + 1$$
, for $i = 2,3,...,n$.

$$f(v_i) = i + 1$$
, for $i = 2,3,...,n$.

Then
$$e_f(0) = e_f(1) = \frac{n^2 - 1}{2}$$
, when n is odd.

$$\text{Then } e_f(0) = \frac{n^2-2}{2} \ \text{ and } e_f(1) = \frac{n^2}{2} \text{ , when } n \text{ even. Thus, } |e_f(0)-e_f(1)| \leq 1.$$

Hence, $(K_n(e)K_n)_{ecc}$ is sdc-graph, for $n \ge 3$.

Illustration: 3.1

K₄(e)K₄ and sdc-labeling of (K₄(e)K₄)_{ecc} are shown in figure 3.1





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Theorem 3.2 $(S(K_{1,n})(e)S(K_{1,n}))_{ecc}$ is sdc-graph for $n \ge 2$.

$$f(w) = 1$$
 and $f(v) = 3n+2$

$$f(w_{1i}) = i+1$$
, for $i = 1,2,...,n$.

$$f(w_{2i}) = i+2n+1$$
, for $i = 1,2,...,n$.

$$f(v_{2i}) = i+n+1$$
, for $i = 1,2,...,n$.

$$f(v_{1i}) = i+3n+2$$
, for $i = 1,2,...,n$.

Then
$$e_f(0)=e_f(1)=\frac{3n^2+2n}{2}$$
 , when n is even and $e_f(0)=\frac{3n^2+2n+1}{2}$ and

$$e_f(1) = \frac{3n^2 + 2n - 1}{2}$$
, when n odd. Thus, $|e_f(0) - e_f(1)| \le 1$.

Thus, $(S(K_{1,n})(e)S(K_{1,n}))_{ecc}$ is sdc-graph for $n \ge 2$.

Illustration: 3.2

 $S(K_{1,3})(e)S(K_{1,3})$ and sdc-labeling of $(S(K_{1,3})(e)S(K_{1,3}))_{ecc}$ are shown in figure 3.2

Theorem 3.3 $(S(K_{1,n})(e)K_m)_{ecc}$ is sdc-graph for $n \ge 2$ and $m \ge 3$.

 $\textbf{Proof:} \ \text{Let} \ G \ \text{be graph} \ S(K_{1,n})(e)K_m. \ G \ \text{has} \ 2n+m+1 \ \text{vertices and} \ 2n+1+\frac{m(m-1)}{2} \ \ \text{edges}.$

Let $(S(K_{1,n})(e)K_m)_{ecc}$ be the eccentric graph of $S((K_{1,n})(e)K_m)$.

The vertices of G are $v,v_{11},v_{12},v_{13},...,v_{1n},v_{21},v_{22},v_{23},...,v_{2n},w_{1},w_{2},w_{3},...,w_{n}$.

Then $K_1 + \overline{K_n} + \overline{K_{m-1}} + \overline{K_{m-1}}$ by adding a vertex to the vertices of first set of vertices in $\overline{K_n}$ and set of vertices in $\overline{K_{m-1}}$, the vertices are $v,v_{11},v_{12},v_{13},...,v_{1n}, v_{21},v_{22},v_{23},...,v_{2n}, w_1,w_2,w_3,...,w_m$. In $(S(K_{1,n})(e)K_m)_{ecc}, w_1$ is mutually adjacent to $v_{21}, v_{22}, v_{23},..., v_{2n}, v_{21}, v_{22}, v_{23},..., v_{2n}$ are mutually to $w_2, w_3,..., w_m$, and also $w_2, w_3,..., w_m$ and $v,v_{11},v_{12},v_{13},...,v_{1n}$ are mutually adjacent. Then, $|V((S(K_{1,n})(e)K_m)_{ecc})| = 2n+m+1$ and $|E((S(K_{1,n})(e)K_m)_{ecc})| = 2nm+m+n-1$.

Define $f: V((S(K_{1,n})(e)K_m)_{ecc}) \rightarrow \{1, 2, ..., 2n+m+1\}$ as follows.

Case 1: n is odd and m is even.

$$f(w_1) = 1$$
 and $f(v) = 2$

$$f(w_i) = i+n+1$$
, for $i = 2,3,...,m$.

$$f(v_{1i}) = i+n+m+1$$
, for $i = 1,2,...,n$.

$$f(v_{2i}) = i+2$$
, for $i = 1,2,...,n$.





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Then
$$e_i(0) = \frac{2nm+m}{2}$$
 and $e_i(1) = \frac{2nm+m-2}{2}$, for n is odd and m is even.

Thus, $|e_f(0) - e_f(1)| \le 1$.

Therefore, $(S(K_{1,n})(e)K_m)_{ecc}$ is sdc graph, for n is odd and m is even.

Case 2: n is odd and m is odd or n is even and m is odd or n is even and m is even.

$$f(w_1) = 1$$
 and $f(v) = 2n+m+1$

$$f(w_i) = i+n$$
, for $i = 2,3,...,m$.

$$f(v_{1i}) = i+n+m$$
, for $i = 1,2,...,n$.

$$f(v_{2i}) = i+1$$
, for $i = 1, 2, ..., n$.

Then
$$e_f(0) = e_f(1) = \frac{2nm+m-1}{2}$$
 , when m is odd and n is either odd or even.

Then
$$e_f(0) = \frac{2nm + m}{2}$$
 and $e_f(1) = \frac{2nm + m - 2}{2}$, when n is even and m is even

Thus,
$$|e_f(0) - e_f(1)| \le 1$$
.

Therefore, $(S(K_{1,n})(e)K_m)_{ecc}$ is sdc-graph for n is odd and m is odd or n is even and m is odd or n is even and m is even. Thus, $(S(K_{1,n})(e)K_m)_{ecc}$ is sdc-graph for $n \ge 2$ and $m \ge 3$.

Illustration: 3.3

 $S(K_{1,3})(e)K_4$ and sdc-labeling of $(S(K_{1,3})(e)K_4)_{ecc}$ sdc-graph are shown figure 3.3.

Theorem 3.4 $(C_n(e)C_n)_{ecc}$ is sdc-graph for $n \ge 3$.

Proof. Let G be $C_n(e)C_n$ graph. G has 2n vertices and 2n+1 edges. The vertices of G are $v_1, v_2, ..., v_n$, $w_1, w_2, ..., w_n$. Let $(C_n(e)C_n)_{ecc}$ be the eccentric graph of $C_n(e)C_n$.

Case 1: when n is even.

Then $(C_n(e)C_n)_{ecc}$ is $S_{n-1,n-1}$, the vertices of G_e are two centers $w_{\frac{n+2}{2}}$, $v_{\frac{n+2}{2}}$ and two set of pendent vertices $v_1, v_2, v_3, \ldots, v_n$

$$v_{\frac{n}{2}}, v_{\frac{n+4}{2}}, \dots, v_n \text{ and } w_1, w_2, w_3, \dots, \quad w_{\frac{n}{2}}, w_{\frac{n+4}{2}}, \dots, w_n. \text{ Then, } |V((C_n(e)C_n)_{ecc})| = 2n \text{ and } |E((C_n(e)C_n)_{ecc})| = 2n-1.$$

Define $f: V((C_n(e)C_n)_{ecc}) \rightarrow \{1, 2, ..., 2n\}$ as follows.

f(
$$v_{\frac{n+2}{2}}$$
) = 1, f(v_1) = 3 and f($w_{\frac{n+2}{2}}$) = 3

$$f(v_i) = i + 2$$
, for $i = 2,3,...,\frac{n}{2}$.

$$f(v_i)=i+1,\, for\,\, i=\frac{n+4}{2}\,,\,\,\frac{n+6}{2},...,n.$$





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$$f(w_i) = i + n + 1$$
, for $i = 1, 2, ..., \frac{n}{2}$.

$$f(w_i) = i + n$$
, for $i = \frac{n+4}{2}$, $\frac{n+6}{2}$,...,n.

Then $e_f(0) = n$ and $e_f(1) = n-1$. Thus, $|e_f(0) - e_f(1)| \le 1$.

Therefore, $(C_n(e)C_n)_{ecc}$ is sdc-graph for n is even.

Case 2: when n is odd.

Then
$$(C_n(e)C_n)_{ecc}$$
 is $\overline{K_{n-2}} + \overline{K_2} + \overline{K_2} + \overline{K_{n-2}}$, the vertices of G are $w_1, w_2, w_3, ..., w_{\frac{n-1}{2}}, w_{\frac{n+5}{2}}, ..., w_n$ of $\overline{K_{n-2}}$,

$$v_{\frac{n+1}{2}}, v_{\frac{n+3}{2}} \text{ of } \overline{K_2}, w_{\frac{n+1}{2}}, w_{\frac{n+3}{2}} \text{ of } \overline{K_2} \text{ and } v_1, v_2, v_3, ..., v_{\frac{n-1}{2}}, v_{\frac{n+5}{2}}, ..., v_n \text{ of } \overline{K_{n-2}}. \text{ Then, } |V((C_n(e)C_n)_{ecc})| = 2n$$

and $|E((C_n(e)C_n)_{ecc})| = 4n-4$.

Define $f: V((C_n(e)C_n)_{ecc}) \rightarrow \{1, 2, ..., 2n\}$ as follows.

f(
$$v_{\frac{n+1}{2}}$$
) = 1 and f($v_{\frac{n+3}{2}}$) = 2, f($w_{\frac{n+1}{2}}$) = 3 and f($w_{\frac{n+3}{2}}$) = 4

$$f(v_i) = n+2+i$$
, for $i = 1,2,..., \frac{n-1}{2}$.

$$f(v_i) = n+i$$
, for $i = \frac{n+5}{2}$, $\frac{n+7}{2}$,...,n.

$$f(w_i) = 4+i$$
, for $i = 1, 2, ..., \frac{n-1}{2}$.

$$f(w_i) = 2+i$$
, for $i = \frac{n+5}{2}$, $\frac{n+6}{2}$,...,n.

Then $e_f(0) = e_f(1) = 2n - 2$. Thus, $|e_f(0) - e_f(1)| \le 1$.

Therefore, $(C_n(e)C_n)_{ecc}$ is sdc-graph for n is odd.

Hence, $(C_n(e)C_n)_{ecc}$ is sdc-graph for $n \ge 3$.

Illustration 3.4. $C_5(e)C_5$ and sdc-labeling of $(C_5(e)C_5)_{ecc}$ are shown in figure 3.4.

CONCLUSIONS

It is confirmed that all of the intriguing and wonderful graph structures of a few more new families of eccentric graphs are sdc-graphs. The sdc-labeling of $(K_n(e)K_n)_{ecc}$, $(S(K_{1,n})(e)S(K_{1,n})_{ecc}$, $(S(K_{1,n})(e)K_m)_{ecc}$ and $(C_n(e)C_n)_{ecc}$ were the only topics we covered.

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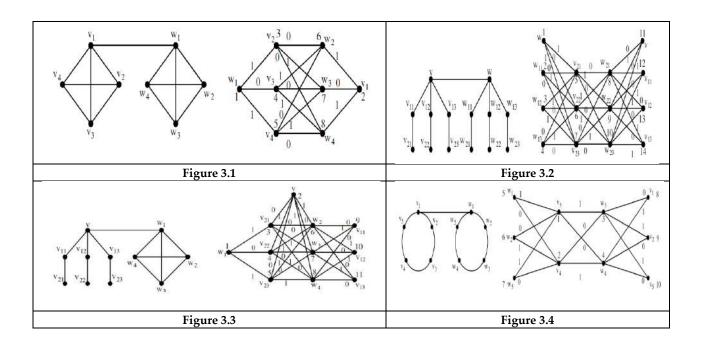
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RESEARCH ARTICLE

Best Proximity Pair Theorem for Generalized Reich type Cyclic Meir-**Keeler Contraction Mapping on Topological Spaces**

S. Sahayarajjoseph Nirmalkumar¹ and S. Sujith²

¹Reg. No: 20221282091006, Research Scholar, PG and Research Department of Mathematics, St.Xavier's College (Autonomous), Palayamkottai, (Affliated to Manonmaniam Sundaranar Abisekapatti-627012, Tirunelveli, Tamil Nadu, India.

²Assistant Professor, PG and Research Department of Mathematics, St. Xavier's College (Autonomous), Palayamkottai, (Affliated to Manonmaniam Sundaranar University), Abisekapatti-627012, Tirunelveli, Tamil Nadu, India.

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*Address for Correspondence

S. Sahayarajjoseph Nirmalkumar

Reg. No: 20221282091006,

Research Scholar,

PG and Research Department of Mathematics,

St.Xavier's College (Autonomous),

Palayamkottai, (Affliated to Manonmaniam Sundaranar University),

Abisekapatti-627012, Tirunelveli, Tamil Nadu, India.

Email: lssnirmal@gmail.com



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ABSTRACT

In this paper, existence and uniqueness of best proximity pairs for generalized Reich type cyclic Meir-Keeler contraction mapping in arbitrary topological space is proved.

Keywords: cyclic contraction, cyclic Meir-Keeler contraction, cyclic topologically contraction, cyclic Reich type contraction, topological P-property.

AMS Subject Classification: 47H09, 47H10.

INTRODUCTION

In [3], Sankar Raj et al., introduced cyclic topologically r-contractive map and proved the existence and uniqueness of best proximity points. Later Sahayarajjoseph Nirmalkumar et al.[4], introduced cyclic topologically contraction map and proved the existence and uniqueness of best proximity points. For other best proximity point result on topological spaces, see[[5],[6]]. In this paper, we have verified the existence and uniqueness of best proximity point for the new class of generalized Reich type cyclic Meir-Keeler contraction maps on topological spaces. The notations below are used throughout the document:





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Consider a topological space Y and a mapping $q: Y \to Y$. Fix $e_0 \in Y$. The orbit $O(e_0, q)$ of q starting at e_0 is defined as $O(e_0, q) := \{q^n(e_0): n \in \mathbb{N} \cup \{0\}\},$

Where $q^0(e_0) = e_0$ and $q^n(e_0) = q(q^{n-1}e_0)$, for all $n \in \mathbb{N}$. Suppose that $\omega: Y \to Y$ is a function. Then the image of $O(e_0,q)$ under ω is $\omega \big(O(e_0,q)\big) = \big\{\omega \big(q^n(e_0)\big) : n \in \mathbb{N} \cup \{0\}\big\}$. The limit point set $Lim\{e_n\}$ of a sequence $\{e_n\}$ in a topological space Y is defined as

$$Lim\{e_n\}\coloneqq\bigcap_{n=1}^\infty\overline{\{e_m\!:\! m\geq n\}},$$

Where \overline{B} denote the closure of B in Y. It is easy to verify that if $q: Y \to Y$ is a continuous mapping and $x \in \text{Lim } O(e_0, q)$, then $\omega(x) \in \text{Lim } \omega(0(e_0, q))$

Main Result

Let us now define the concept of generalized Reich type cyclic Meir-Keeler contraction mapping [1] on a topological space

Definition 2.1. Let E and F be any two non-empty subsets of a topological space Y. A cyclic mapping $q: E \cup F \to E \cup F$ is called generalized Reich type cyclic topologically Meir-Keeler contraction map w.r.t a continuous function $\omega: Y \times Y \to \mathbb{R}$ for some $u \in E$ and for each $\varepsilon > 0$,

there exists a $\delta > 0$ such that

$$R(u,v) < D_{\omega}(E,F) + \epsilon + \delta \quad \text{implies} \mid \omega(qu,qv) \mid < D_{\omega}(E,F) + \epsilon, \forall u \in E, \forall v \in F, \tag{2.1}$$

Where (u, v) = $\frac{1}{3}$ [| ω (u, v) | +| ω (qu, u) | +| ω (qv, v) |] .

Here's an example

Example 2.2. Consider \mathbb{R}^2 with its standard topology. Let $E := \{(0,t): 0 \le t \le 1\}$ and $F := \{(1,s): 0 \le s \le 1\}$. Let $q: E \cup F \to E \cup F$ be defined as

$$q(u,v) \coloneqq \begin{cases} \left(1,\frac{v}{2}\right), & \text{if } u = 0, \\ \left(0,\frac{v}{2}\right), & \text{if } u = 1. \end{cases}$$

Let $\omega: \mathbb{R}^2 \times \mathbb{R}^2 \to \mathbb{R}$ be a mapping defined as $\omega((u, v), (w, z)) = vz$ for all $(u, v), (w, z) \in \mathbb{R}^2$.

 ω is obviously a continuous function, not a metric on \mathbb{R}^2 . Observe that $D_{\omega}(E,F)=0$.

If for $u \in E$ and $v \in F$, $|\omega(u,v)| > D_{\omega}(E,F)$ then take $\epsilon = |\omega(u,v)| - D_{\omega}(E,F)$. It is easy to verify that for this $\epsilon > 0$ there exists $\delta > R(u,v)$ such that the function q satisfies the conditions of generalized Reich type cyclic topologically Meir-Keeler contraction map w.r.t ω .

Let us now show that the best proximity point in a topological space exists and is unique:

Proposition 2.3. Let E and F be non-empty and closed subsets of topological space Y. Let

 $q: E \cup F \to E \cup F$ be a generalized Reich type cyclic topologically Meir-Keeler contraction map w.r.t a continous function $\omega: Y \times Y \to \mathbb{R}$. If $u \in E$ satisfies condition (2.1), then $|\omega(q^n u, q^{n+1}u)| \to D_{\omega}(E, F)$ as $n \to \infty$.

Proof. Suppose q is generalized Reich type cyclic topologically Meir-Keeler contraction map w.r.t a continuous function ω . Take $u \in E$ for which (2.1) holds.

Since either n or n + 1 is even, then for each $u \in E$, we have,

$$\frac{1}{3}[|\omega(q^n u, q^{n-1}u)| + |\omega(q^{n+1}u, q^n u)| + |\omega(q^n u, q^{n-1}u)|] \ge D_{\omega}(E, F).$$

Consider the case, $\frac{1}{3}[|\omega(q^{n}u, q^{n-1}u)| + |\omega(q^{n+1}u, q^{n}u)| + |\omega(q^{n}u, q^{n-1}u)|] = D_{\omega}(E, F).$

Then due to (2.1), we have $|\omega(q^{n+1}u, q^nu)| < D_{\omega}(E, F) + \epsilon$,

Which is equivalent to



(2.2)



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 $\mid \omega(q^{n+1}u,q^nu) \mid < \frac{1}{3} \left[\mid \omega(q^nu,q^{n-1}u) \mid + \mid \omega(q^{n+1}u,q^nu) \mid + \mid \omega(q^nu,q^{n-1}u) \mid \right] + \epsilon.$

Thus we have, $|\omega(q^{n+1}u, q^nu)| \le |\omega(q^nu, q^{n-1}u)|$, as $\epsilon \to 0$.

Now consider the other case, that is

$$\begin{split} \frac{1}{3} [\mid \omega(\mathbf{q}^{\mathbf{n}}\mathbf{u}, \mathbf{q}^{\mathbf{n}-1}\mathbf{u} \mid + \mid \omega(\mathbf{q}^{\mathbf{n}+1}\mathbf{u}, \mathbf{q}^{\mathbf{n}}\mathbf{u}) \mid + \mid \omega(\mathbf{q}^{\mathbf{n}}\mathbf{u}, \mathbf{q}^{\mathbf{n}-1}\mathbf{u}) \mid] > D_{\omega}(E, F). \\ \text{Set } \epsilon_{1} &= \frac{1}{3} [\mid \omega(\mathbf{q}^{\mathbf{n}}\mathbf{u}, \mathbf{q}^{\mathbf{n}-1}\mathbf{u}) \mid + \mid \omega(\mathbf{q}^{\mathbf{n}+1}\mathbf{u}, \mathbf{q}^{\mathbf{n}}\mathbf{u}) \mid + \mid \omega(\mathbf{q}^{\mathbf{n}}\mathbf{u}, \mathbf{q}^{\mathbf{n}-1}\mathbf{u}) \mid] - D_{\omega}(E, F) > 0. \end{split}$$

According to (2.1), for this ϵ_1 , there exists δ_1 such that

$$|\omega(q^{n+1}u,q^nu)| < D_{\omega}(E,F) + \epsilon_1 = \frac{1}{3}[|\omega(q^nu,q^{n-1}u)| + |\omega(q^{n+1}u,q^nu)| + |\omega(q^nu,q^{n-1}u)|].$$

Hence, $|\omega(q^{n+1}u, q^nu)| \le |\omega(q^nu, q^{n-1}u)|$ for all $n \in \mathbb{N}$.

Let $\alpha_n = |\omega(q^{n+1}u, q^nu)|$. Clearly $\{\alpha_n\}$ is a decreasing sequence bounded below by $D_{\omega}(E, F)$.

Therefore $\{\alpha_n\}$ converges to some α with $\alpha \geq D_{\omega}(E, F)$.

We now show that $\alpha = D_{\omega}(E, F)$ by assuming the contrary, that is $\alpha > D_{\omega}(E, F)$.

Set $\epsilon = \alpha - D_{\omega}(E,F) > 0$. Then there exists $\delta > 0$ for which (2.1) holds. Since $\{ |\omega(q^{n+1}u,q^nu)| \} \rightarrow \alpha$, there exist a $n_0 \in \mathbb{N}$ such that

$$\alpha \leq \frac{1}{3}[\mid \omega(\mathsf{q}^n\mathsf{u},\mathsf{q}^{n-1}\mathsf{u})\mid +\mid \omega(\mathsf{q}^{n+2}\mathsf{u},\mathsf{q}^{n+1}\mathsf{u})\mid +\mid \omega(\mathsf{q}^{n+1}\mathsf{u},\mathsf{q}^n\mathsf{u})\mid] < \epsilon + D_{\omega}(E,F) + \delta, \forall n \geq n_0.$$
 Thus, $\mid \omega(\mathsf{q}^{n+2}\mathsf{u},\mathsf{q}^{n+1}\mathsf{u})\mid < D_{\omega}(E,F) + \epsilon = \alpha, \ \forall n \geq n_0.$ Which is contradiction .

Hence $\alpha = D_{\omega}(E, F)$.

Theorem 2.4. Let E, F be nonempty and closed subsets of a topological space Y and $\omega: Y \times Y \to R$ be a continuous mapping. Let q: E∪F → E∪F be a continuous generalized Reich type cyclic topologically Meir-Keeler contraction mapping w.r.t ω . Then for any $x \in E \cup F$, the set Lim O(x,q) is either empty or consist of a point {u}, which satisfy $|\omega(u, qu)| = D_{\omega}(E, F)$.

Proof. Let $x \in E \cup F$. Let's say Lim O(x, q) is nonempty, by continuity of q if $v \in Lim O(x, q)$,

$$q(v) \in \text{Lim } q(O(x,q)) \subseteq \text{Lim } O(x,q).$$

Define $\tau : E \cup F \to R$ by $\tau (u) := |\omega(u, qu)|$. Since q, ω are continuous, τ is also continuous. Hence,

 $\tau(v) \in \text{Lim } \tau(O(x,q))$

Suppose that $|\omega(q^nx,q^{n+1}x)| \neq D_{\omega}(E,F)$ for all $n \in \mathbb{N}$. Since q is generalized Reich type cyclic topologically Meir-Keeler contraction map w.r.t ω , from Proposition 2.3, we have $\{\tau(q^nx)\}\to D_\omega(E,F)$ asn $\to\infty$. i.e., Lim $\tau(O(x,q))$ is a singleton set. From (2.3), $\tau(v) = D_{\omega}(E, F)$.i.e., $|\omega(v, qv)| = D_{\omega}(E, F)$.i.e., v is a best proximity point q with respective ω . Suppose that $|\omega(q^{n_0} x, q^{n_0+1}x)| \neq D_{\omega}(E, F)$ for some $n_0 \in \mathbb{N}$. Then $\{\tau(q^n x)\}$ becomes an eventually constant sequence with the constant value $D_{\omega}(E,F)$. Hence Lim $\tau(O(x,q))$ is a singleton set. The same conclusion follows by the preceding argument.

Theorem 2.5. Let E, F be nonempty and closed subsets of a topological space Y and $\omega: Y \times Y \to R$ be a continuous mapping that satisfies the following requirements:

(1)
$$\omega(u, v) = \omega(v, u)$$
, for all $u \in E, v \in F$,

(2)
$$\omega(u_1, u_2) = 0 \iff u_1 = u_2$$
, for all $u_1, u_2 \in E$ (or $u_1, u_2 \in F$).

Let q: E ∪ F → E ∪ F be a continuous generalized Reich type cyclic topologically Meir-Keeler contraction mapping w.r.t ω . Suppose that the pair (E, F) has P-property w.r.t ω . Then for any $x \in E \cup F$, the set Lim O(x,q) is either empty or Lim $O(x,q) \cap E$ is a singleton set $\{u\}$, which satisfy $|\omega(u,qu)| = D_{\omega}(E,F)$.

Proof. By Theorem 2.4, we conclude that the set Lim O(x,q) is either empty or contains a best proximity point v(say)of q w.r.t ω . Now we show that the best proximity point, v, is unique. Without loss of generality, assume that v is in E. Suppose there exists two points u_0, v_0 in E such that $u_0, v_0 \in \text{Lim } O(x, q)$ satisfying

$$|\omega(\mathbf{u}_0, \mathbf{q}\mathbf{u}_0)| = D_{\omega}(E, F) \tag{2.4}$$

$$|\omega(\mathbf{v}_0, \mathbf{q}\mathbf{v}_0)| = D_{\omega}(E, F) \tag{2.5}$$





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Since q is generalized Reich type cyclic topologically Meir-Keeler contraction mapping w.r.t ω and by Proposition 2.3, we have, $|\omega(qu_0, q^2u_0)| = D_{\omega}(E, F)$.

Hence, $|\omega(u_0, qu_0)| = |\omega(q^2u_0, qu_0)| = D_{\omega}(E, F)$. By P-property, we have $|\omega(u_0, q^2u_0)| = |\omega(qu_0, qu_0)| = 0$. Thus $u_0 = q^2u_0$. Similarly $v_0 = q^2v_0$.

Now by equation (2.2), we have,

$$|\omega(\mathbf{v}_0, \mathbf{q}\mathbf{u}_0)| = |\omega(\mathbf{q}^2\mathbf{v}_0, \mathbf{q}\mathbf{u}_0)|$$

$$\leq |\omega(q\mathbf{v}_0, \mathbf{u}_0)|. \tag{2.6}$$

Similarly,
$$|\omega(qv_0, u_0)| \le |\omega(v_0, qv_0)|$$
 (2.7)

From (2.6) and (2.7), we get that

$$|\omega(v_0, qu_0)| = |\omega(u_0, qv_0)|. \tag{2.8}$$

Suppose that $|\omega(qu_0, v_0)| > D_{\omega}(E, F)$. Then $\epsilon_0 = |\omega(qu_0, v_0)| - D_{\omega}(E, F) > 0$.

Claim: For this ϵ_0 , there exists a $\delta_0 > 0$ satisfying $R(qu_0, v_0) < D_{\omega}(E, F) + \epsilon_0 + \delta_0$.

Assume the contrary, Suppose for all $\delta_0 > 0$, $R(qu_0, v_0) \ge D_{\omega}(E, F) + \epsilon_0 + \delta_0$.

That is,

$$\begin{split} 1/\,3\,[|\omega(\mathsf{qu}_0,\mathsf{v}_0)| \,+\, |\omega(\mathsf{q}^2\mathsf{u}_0,\mathsf{qu}_0)| \,+\, |\omega(\mathsf{qv}_0,\mathsf{v}_0)|] \,\geq\, D_\omega(E,F) \,+\, \epsilon_0 \,+\, \delta_0 \\ 1/\,3\,[|\omega(\mathsf{qu}_0,\mathsf{v}_0)| \,+\, 2D_\omega(E,F)] \,\geq\, D_\omega(E,F) \,+\, \epsilon_0 \,+\, \delta_0 \\ 1/\,3\,|\omega(\mathsf{qu}_0,\mathsf{v}_0)| \,\geq\, (\,1\,-\,2/\,3\,)D_\omega(E,F) \,+\, \epsilon_0 \,+\, \delta_0 \\ |\omega\left(\mathsf{qu}_0,\mathsf{v}_0\right)| \,\geq\, D_\omega(E,F) \,+\, 3\, \epsilon_0 \,+\, 3\, \delta_0 \\ |\omega(\mathsf{qu}_0,\mathsf{v}_0)| \,\geq\, D_\omega(E,F) \,+\, 3\, (|\omega(\mathsf{qu}_0,\mathsf{v}_0)| \,-\, D_\omega(E,F) \,+\, 3\, \delta_0 \\ |\omega(\mathsf{qu}_0,\mathsf{v}_0)| \,\leq\, D_\omega(E,F) \,+\, 3\, \delta_0 \\ |\omega(\mathsf{qu}_0,\mathsf{v}_0)| \,\leq\, D_\omega(E,F) \,-\, 3/\, 2\, \delta_0 \end{split}$$

As $\delta_0 \to 0$, $|\omega(qu_0, v_0)| < D_{\omega}(E, F)$ a contradiction.

Therefore, $R(qu_0, v_0) < D_{\omega}(E, F) + \epsilon_0 + \delta_0$. Then due to (2.1), we have,

$$\begin{aligned} |\omega (\mathsf{q}^2 \mathsf{u}_0, \mathsf{q} \mathsf{v}_0)| &< D_{\omega}(E, F) + \epsilon_0 \\ &= D_{\omega}(E, F) + |\omega (\mathsf{q} \mathsf{u}_0, \mathsf{v}_0)| - D_{\omega}(E, F) \\ &= |\omega (\mathsf{q} \mathsf{u}_0, \mathsf{v}_0)|. \end{aligned}$$

That is, $|\omega(u_0, qv_0)| < |\omega(qu_0, v_0)|$, which is a contradiction to the equation (2.8).

Therefore, $|\omega(qu_0, v_0)| = D_{\omega}(E, F)$.

By using equation (2.4) and P-property, $|\omega(u_0, qu_0)| = |\omega(v_0, qu_0)| = D_{\omega}(E, F)$

 $\Rightarrow |\omega(u_0, v_0)| = |\omega(qu_0, qu_0)| = 0$. Hence $u_0 = v_0$. i.e., the best proximity point of q w.r.t ω is unique in E.

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RESEARCH ARTICLE

Analysis on Online Learning of Mathematics and Student's Attitude

A. Aruna¹ and PL. Meenakshi²

¹Post Graduate Student, Department of Mathematics, Avinashilingam Institute for Home Science and Higher Education for Women, Tamil Nadu, India.

²Assistant Professor, Department of Mathematics, Avinashilingam Institute for Home Science and Higher Education for Women, Tamil Nadu, India.

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*Address for Correspondence

A. Aruna

Post Graduate Student, Department of Mathematics, Avinashilingam Institute for Home Science and Higher Education for Women, Tamil Nadu, India.



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ABSTRACT

The COVID-19 pandemic has resulted in significant effects in various sectors, digitalising the world. One of the most impacted sectors includes education, which resulted in the widespread adoption of remote learning policies like video classes, system-proctored tests etc. However, the shift to online classes has made learning more challenging for students as the accessibility to smartphones and systems is not even. It made understanding Mathematics even more challenging as it requires effective student-teacher interaction to understand the subject and cope with Mathematics and fewer students are enrolling in this field. This study aims to explain why students' interest in enrolling in Undergraduate Mathematics Courses in Coimbatore district has decreased after COVID-19. It continues to investigate whether the 11th and 12th graders' interest in Mathematics has been impacted by their online math education. Also, the relationship between online learning of Mathematics and students' attitudes towards math is analysed.

Keywords: Online learning, COVID-19, Attitude, t-test.

INTRODUCTION

Mathematics is widely known as the "language of the universe" and includes a wide range of ideas, hypotheses, and applications that are essential to many academic disciplines as well as in daily life. Mathematics is the foundation of every innovation. It is widely used in almost every field. Its wide applications are seen in technologies, software, computers, art, economy, engineering, sports, and even in contemporary and ancient architectures. From basic algebra to sophisticated calculus, geometry to simple arithmetic, and all points in between, Mathematics offers the foundation for comprehending quantitative connections, resolving issues, and making wise judgments. Mathematics is a crucial subject in school and college curricula. It is a compulsory subject at the school level and a major subject at





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a higher level. However, many students find Mathematics challenging, resulting in many of them failing in this subject. Despite the importance of Mathematics in research and development it does not attract many of the students to take it as a major. Most students find Mathematics difficult and online learning during COVID-19 made it a more and more difficult subject. It is found that in the aftermath of the pandemic enrolment of students in Mathematics is largely impacted negatively. This study aims to identify the online learning on student's attitudes towards Mathematics.

ONLINE LEARNING

Electronic devices like mobile, laptops, computers, etc.., are used for learning, a sort of educational activity with the support of the internet facilitating student-teacher interactions and instruction. Education around the world has been impacted by the development of online learning brought about by COVID-19. There will always be a process of interaction in every learning action, whether it takes place online or in person. The three types of interaction in the learning process are the interaction between

- 1. students and students,
- 2. students with teachers and
- 3. students with the material (Bernard et al., 2009).

Mathematicsis among the subjects that are affected by online learning. Here are some aspects that need to be considered regarding online learning of Mathematics during the pandemic and the student's attitudes towards the subject. The aspects are:

- 1. Challenges of Online Learning in Mathematics
- 2. Adaptation to Online Platforms
- 3. Students' attitude towards Mathematics
- 4. Long-term impact.

STUDENTS' ATTITUDE TOWARDS MATHEMATICS:

The term "attitude" describes a person's recurring beliefs, emotions, and actions that reveal their character and point of view. It is an amalgam of attitude, conviction, feeling, and action that mould reactions to different circumstances. A person's attitude greatly affects how they handle problems, relates to other people, and make decisions. Attitudes can be neutral, negative, or positive. It is essential for influencing general well-being, directing behaviours, and forming perceptions. A person's attitude is dynamic and subject to change in response to events, cultural standards, and personal development. To build healthy connections and succeed both personally and professionally, it is crucial to recognize and control one's attitude. The approach that students take to mathematics has a significant impact on both their academic achievement and interest in the subject. Students who take enjoyment in performing mathrelated assignments demonstrate a good attitude toward the subject. Therefore, it is essential to maintain a favourable attitude towards Mathematics to promote students' willingness to learn the subject and include it in Mathematics instruction. Conversely, a negative attitude towards Mathematics can lead to reluctance to learn and spend time on the subject, thus hindering their progress. Factors such as inappropriate instructional materials, poor instructional practices, classroom management, and class activities can significantly affect students' attitudes towards Mathematics.

REVIEW OF LITERATURE

Lazaros Anastasiadis and Poulcheria Zirinoglou (2022)

The study's goal is to determine how Greek students feel about studying analytics, AI algorithms, and big data. Four conceptual constructs are used in the study to examine the attitudes of the students: motivation, enjoyment, value, and self-confidence. The relationship between value and enjoyment, drive and value, and self-confidence and value were shown to have the greatest connections. Furthermore, the findings demonstrated that there was no big difference between male and female students' views about Mathematics.





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I Putu Wisna Ariawan (2022)

The study demonstrated the effects of COVID-19 on the academic performance of Singaraja City public high school students enrolled in online mathematics courses. Some scams within the area of virtual math classes misled learners into thinking that their academic achievements were poor and that their abilities were compared to those of another. The results indicate that to eliminate the negative impact on learning teachers should put more effortinto neutralizing the effect in upcoming academic years.

Justice Enu, Osei K Agyman, Daniel Nkum (2015)

The purpose of the study was to investigate the factors impacting the mathematics performance of college students. The study concluded that the student's performance in mathematics is impacted by some issues like the style of instruction, which makes the student a passive participant in the learning process, and inadequate teaching and learning resources.

Nazir Haider Shah, Nadia Nazir (2023)

According to the study, anxiety had a higher mean score when compared to motivation, confidence, and importance in terms of students' attitudes towards mathematics. It indicates that students have anxiety before beginning a math lesson. It was determined that secondary school male and female students did not significantly differ in their mathematical performance and that male students had a greater attitude toward mathematics than female students. Anxiety, significance, getting around, and confidence all have a significant impact on mathematics success. The parameters of students' attitudes indicate that they will not change in the future with regard to their interest in mathematics; however, confidence and motivation will play a larger role than significance, and anxiety it will prevent students from interested in learning challenging subjects like mathematics.

THE PURPOSE OF THE STUDY

This study aims to explain why students' interest in enrolling in Undergraduate Mathematics Courses in Coimbatore district has decreased after COVID-19. It continues to investigate whether the 11th and 12th graders' interest in Mathematics has been impacted by their online math education. Also, the relationship between online learning of Mathematics and students' attitudes towards math is analysed.

LIMITATIONS OF THE STUDY

The focus of the study is limited to female undergraduate students of educational institutes who are not pursuing Mathematics as their major subject. The nature of the study is descriptive and qualitative, which means that it aims to provide an in-depth understanding of the research topic.

RESEARCH METHODOLOGY

An educational institute was selected for data collection, and the study population targeted ordinary-level students. As a result, the researchers randomly selected 50 students who participated in the study and had an average age of 18 years. To address the research question, a descriptive survey design was employed in the present study. The researchers constructed questionnaires based on the research questions, which were divided into two main themes:

- 1. Online learning of Mathematics, and
- 2. Students' attitude towards Mathematics.

The researchers deemed six items that are sufficient to analyse the relationship between online learning of Mathematics and students' attitudes. The students were given the surveys in English, using a 4-point Likert scale ranging from 4 (strongly agree) to 1 (strongly disagree).

DATA ANALYSIS

Data were analyzed using Microsoft Excel. The relation between students' attitudes toward mathematics and their online mathematics education was examined utilizing





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- 1. independent sample t-tests.
- 2. Mean
- 3. Standard deviation

was utilized to describe the Online learning of Mathematics and student attitude.

Note: df – degree of freedom.

STUDENT'S ATTITUDE TOWARDS MATHEMATICS ONLINE LEARNING MATHEMATICS

RESULT AND FINDINGS

To test the hypotheses, a t-test for independence with a significance level α = 0.05 is used.

Null hypothesis H₀: There is no relationship between online learning and students' attitudes towards Mathematics. Alternative hypothesis H₁: There is a relationship between online learning and students' attitudes towards Mathematics. By comparing the t-value and critical value 4.33277>1.9845So, we accept H₁.This study aimed to explore the correlation between online learning of Mathematics and students' attitudes towards the subject. The study proposed that online learning could impact attitudes towards math. The results show that there is, in fact, a link between students' attitudes toward mathematics and online learning. Table 1 displays the results of the Attitude Towards Mathematics Inventory as rated by the student-respondents. The table includes the mean score for each item and the overall mean rating.

- 1. The highest recorded mean score of 3.34 was achieved by the positive statement "I want to develop my mathematical skills",
- 2. The lowest recorded mean score of 1.74 was obtained by the negative statement "Mathematics is not necessary for our studies".
- 3. The overall mean Attitude towards Mathematics for all students is 2.7033.

Students find the subject something confusing and complicated. Despite these perceptions, they still see Math as important in our studies. Students see Mathematics in everyday living and it's important to everyone. The level of Online learning of Mathematics is presented in Table 2. Here the Highest mean value of 2.88 was obtained by the statement of the items, "Even if I have no idea about the subject matter, I am too lazy to approach the teacher when the math lessons are done online" and "Sometimes, internet network issues prevent me from properly participating in online math periods". Table 3 shows that Online learning of Mathematics is related to student attitude. The relationship between Online learning of Mathematics and the student's attitude was measured using an independent sample t-test. According to the table,

The score of Online Learning is,

N=50,M=2.7033,SD=0.10077

and the score of the student's attitude is,

N=50, M=2.7933, SD=0.10686

Whereas,

t=4.33277>1.9845

So, there was a relationship between Online learning of Mathematics and the student's attitude. Furthermore, the Null hypothesis was rejected.





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CONCLUSION

The purpose of the study was to investigate the impact of online learning on students' attitudes towards Mathematics during the pandemic. The results of the study showed a connection between online learning and students' negative outlook on mathematics. Consequently, online learning has affected students' attitudes and resulted in a low enrollment rate in Mathematics. So, the enrollment rate may rise if we reduce these explanations.

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Table 1.Mean and standard deviation of student's responses in attitude towards Mathematics

Questions	Total	Strongly agree	Agree	Disagree	Strongly disagree	Mean	Standard deviation
Mathematics is very interesting to me and I enjoy Mathematics	50	13	27	9	1	3.04	0.727
courses.	30	13	27	9	1	3.04	0.727
My mind goes blank and I am							
unable to think when working	50	27	6	16	1	2.76	0.687
on Mathematics.	30	27	O	10	1	2.70	0.007
I love the study of Mathematics.							





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	50	11	23	14	2	2.86	0.808
Mathematics is not necessary in our studies	50	1	5	24	20	1.74	0.723
I've never been good at Mathematics, and it's my least favourite subject.	50	7	16	21	6	2.48	0.886
I want to develop my mathematical skills.	50	20	27	3	0	3.34	0.593
					Overall Mean and SD	2.7033	0.10077

Table 2: Mean and standard deviation of student responses in Online learning of Mathematics.

Table 2: Mean and standard deviation	on or st		111562 111	Omme ie		lamema	Standard
Questions	Total	Strongly	Agree	Disagree	Strongly	Mean	
		agree	Ü	Ü	disagree		deviation
I believe that my online learning math							
scores don't accurately represent my							
actual mathematics skills.	50	5	33	12	0	2.86	0.572
Even if I have no idea about the subject							
teacher when the math lessons are					2.88 2.88 2 2 2.88 6 2 2.72 3 3 2.82		
	to idea about the subject to lazy to approach the in the math lessons are one online. 50 7 31 Therefore the math lessons are one online. 50 7 31 Therefore the math lessons are one online. 50 7 31 Therefore the math lessons are one online. 50 7 31 Therefore the math lessons are one online. 50 6 26 The mathematics lessons are one online. 50 7 31	11	1		0.659		
done offine.							
Sometimes, internet network issues							
prevent me from properly participating							
in online math periods.	50	10	26	12	2	2.88	0.773
Math skills have decreased due to							
studying online.	50	6	26	16	2	2.72	0.730
Even though the mathematics lessons							
are done online, I frequently receive	50	10	24	13	3	2 82	0.825
high marks.	50	10	21	10		2.02	0.020
Online math classes have potential							
sufficient opportunities for interactive							
learning and engagement.	50	7	20	19	4	2.60	0.833
				Overall	Mean and		
				Overall	SD	2.7933	0.10686

Table 3 Analysis of Online Learning and Student's Attitude towards Mathematics.

	N	Mean	Standard deviation	df	t-value	Critical value	The result
Online learning	50	2.7033	0.10077				
				98	4.33277	1.9845	Reject Ho
Attitudes	50	2.7933	0.10686				





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RESEARCH ARTICLE

A Diagnosis of Renal Disease using Multiclass with ERF Classifier **Learning Techniques**

K. Kottaisamy^{1*} and S. Chidambaranathan²

¹Research Scholar, Regno. 19221282291017, Dept. of Computer Applications, St. Xavier's College (Autonomous), Palayamkottai, (Affiliated to Manonmaniam Sundaranar University) Tirunelveli, Tamil Nadu, India.

²Associate Professor, Dept. of Computer Applications, St. Xavier's College (Autonomous), Palayamkottai (Affiliated to Manonmaniam Sundaranar University) Tirunelveli, Tamil Nadu, India.

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*Address for Correspondence

K. Kottaisamy

Research Scholar,

Regno. 19221282291017,

Dept. of Computer Applications,

St. Xavier's College (Autonomous), Palayamkottai,

(Affiliated to Manonmaniam Sundaranar University)

Tirunelveli, Tamil Nadu, India. Email: kskottaisamy@gmail.com



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ABSTRACT

Machine learning-based medical diagnostics will make early renal disease stage prediction viable in underdeveloped and developing nations. Multiclass classification algorithms are used to diagnose kidney illness using a real-time dataset. As a result, kidney failure sets in, making an early prediction of Chronic Renal Disease (CRD) vital even if it may be difficult. People with diabetes, hypertension, autoimmune diseases, or a family history of the condition should have priority screening for an early diagnosis of CRD. The suggested Ensemble Random Forest (ERF) with a multiclass classifier model was trained and tested in the study's research using the renal disease dataset. Finally, we considered how accurate the proposed model was at predicting the future. ERF with multi-class classification is used to classify data with 26 features, 20 features, 15 features, and 10 features to predict the Chronic renal stages. Finally, among the 26 characteristics, for the multiclass categorization. For all ten characteristics combined, the accuracy of the renal stage is 97%.

Keywords: Chronic Renal Disease, Diagnosis, Prediction, Multiclass Classification, Machine Learning, Hypertension, Diabetes.





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INTRODUCTION

Biosciences have advanced to a greater level, generating vast amounts of data via electronic health records. This has created an urgent demand for knowledge generation from this massive volume of data. In this branch of biology, machine learning and data mining are crucial. Damaged kidneys that are unable to filter blood as well as they should are the outcome of CRD. People with lower socio-economic positions bear a disproportionate amount of the burden of CRD because they have a higher prevalence of the disease, less access to treatment, and worse outcomes. Therefore, it is crucial for equality that CRD is detected early and treated. CRD can be decreased or even prevented by doing tests such as urine tests, blood tests, and kidney scans, and consulting a doctor about additional symptoms of renal disease. The kidneys do not normally fail all at once in chronic renal disease. Instead, renal disease frequently advances slowly over time. If detected early, medications and lifestyle adjustments may help delay or prevent the course of CKD. CKD has five phases. The five phases of kidney disease are defined by the National Kidney Foundation (NKF). This enables clinicians to offer the best care possible because each stage necessitates a distinct set of tests and treatments. To ascertain the stage of renal illness, physicians utilize the Estimated Glomerular Filtration Rate (eGFR), a mathematical formula that takes into account an individual's age, gender, and serum creatinine level (ascertained by a blood test). An essential biomarker of renal function is creatinine, a byproduct of muscle activity. Blood creatinine levels grow when renal function decreases, whereas when the kidneys are functioning normally, they remove creatinine from the blood. The first stage of renal disease is linked to a high or normal GFR (GFR > 90 mL/min); the second stage is linked to CKD (GFR = 60-89 mL/min); the third stage is linked to moderate CKD (GFR = 45–59 mL/min); the fourth stage is linked to severe CKD (GFR 15 mL/min); and the fifth stage is linked to end-stage CKD.In this study, we created machine learning models utilizing important pathological categories that were specifically chosen to uncover clinical test characteristics that would help with the early and accurate diagnosis of CKD. Time and money will be saved for diagnostic screening with this method.

we investigated the accuracy rate of feature selection approaches in this work by lowering the dimensionality of the feature and merging heterogeneous classifiers to produce the ensemble model. The data employed in this study's response parameter are the various stages of chronic kidney disease based on eGFR, a variable that is categorical with five stages of development (i.e., 1, 2, 3A, 3B, 4, and 5). Several response rate categories bias the data for the above parameter in a negative direction. This work uses statistical models that capture the characteristics of the ordinal response factor to determine the factors impacting GFR to discover the optimum model that more precisely categorizes very unbalanced data. Our findings imply that when employing our suggested machine learning models to diagnose CKD, optimized datasets with key features perform well. Additionally, we assessed the characteristics of clinical tests using blood and urine tests as well as clinical indicators with low acquisition prices. The best-performing prediction model, the random forest classifier, produced excellent levels of CKD diagnosis accuracy using the optimized and pathologically categorized attribute set.

Conclusions: Using machine learning, we were able to produce effective predictive analytics for CKD screening that can be used to improve CKD screening processes and enable more effective and prompt treatment strategies. This document has the following structure. Section 2 demonstrates multiple machine learning Classifier methods for predicting chronic renal disease. Response variables, explanatory factors, and their characteristics are all part of the models and data for this study. The dataset is introduced and explained in Section 3 along with two original methods: preprocessing and model building. The results of a classification performance comparison between ensemble random forest machine learning techniques and generalized multi-class models are examined in Section 4. Finally, Section 5 offers a study analysis. Previously, multiple researchers employed chronic renal illness to test the precision of numerous machine-learning algorithms using methodologies. In their study, seven classifier algorithms were utilized by Chittora et al. [1]: the Chi²automatic interaction detector, Random Forest, ANN, C5.0, Logistic Regression (LR), Linear Support Vector classifier (LSVM) concluded penalty L1 & through penalty L2. The approach that produced the best accuracy was the linear model, which they determined after measuring the correctness, precision, recall, and AUC curve. The results have been calculated independently for each classifier using the least





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absolute shrinking and selection operator regression, synthetic minority oversampling with complete attributes, Wrapper approach feature selection, and full features. According to this study, among all the classifiers, LSVM had the highest accuracy across all experiments. However, the aims for Logistics and KNN were not sufficiently met. Using pre-processing methods and datasets from the UCI Machine Learning repository such as label encoding and min-max normalizing, Sathyaraj et al. [2] presented a hybrid model that can accomplish better to detect kidney disease. Finally, they discovered that the Random Forest and AdaBoost algorithms consistently preserved a higher perfection rate. Wang et al.'s [3] regression model took into account six variables, including sex, age, smoking habit, haemoglobin, and urine protein level, to predict the risk of CKD. They tested with the XGBoost, Random Forest, and ResNet neural network regression models. Finally, XGBoost achieved the greatest result with an R2 score of 0.5523. To raise public awareness, Shirahatti et al. [4] proposed a classification system that uses ML algorithms to evaluate the likelihood of being impacted by CKD. Electronic Medical Records (EMR), Clinical history, and laboratory testing were used to gather the necessary data. They compared some machine learning methods, both with and without preprocessing, and determined that the random forest classifier achieved 100% accuracy. In underdeveloped and emerging nations, early detection of CKDwill be possible because of machine learning-based medical diagnostics. Renal disease is diagnosed with machine learning models using a real-time dataset. As a result, kidney failure sets in, and although early detection of CKD may be difficult, it is crucial. Prioritizing early detection of CKD requires screening those with diabetes, hypertension, autoimmune diseases, or a family history of the illness.

Machine learning may be applied to inpatient electronic medical data to find patterns associated with certain diseases and alert physicians to any abnormalities. The research findings indicate that enhancing the effectiveness of classification algorithms by feature selection can lead to a rise in the precision of chronic kidney disease prediction and diagnosis machine learning approaches including decision tree, random forest classifier, and extra tree. The trial's conclusions could help the medical field anticipate problems and make decisions early on to treat kidney illness and save lives. The experimental results for the proposed model appear to confirm the predicted degree of prediction accuracy [5]. Using the data from 551 patients, Xiao et al. [6] provided an approach to managing 9 machine learning classifiers: KNN, LR, Ridge regression, Random Forest ,Lasso regression, XGB, SVC, and Elastic Net. They applied machine learning, neural networks, and statistical approaches to analysis a variety of demographic information, including 13 blood and 5 additional indicators, to determine the disease's current stage. As a consequence, the suggested approach R-ACO-SVM achieved 99.50% accuracy and obtained good results when compared to the supplementary feature selection methods. Despite the significant number of missing values in the University of California Irvine (UCI) machine learning repository, Chen et al.'s [7] approach for diagnosing CKD makes use of machine learning techniques. They used feed-forward neural networks, random forest, naive Bayes classifiers, support vector classifiers, k-nearest neighbours, and logistic regression. They conducted their suggested model through about 10 simulations, and they discovered that logistic regression and random forest both had an accuracy of 99.83% on average. A patient's CKD status may be ascertained using machine learning algorithms. The dataset consists of 400 records of Indian patients from the Apollo Hospital in Tamil Nadu, along with 25 covariates. The dataset is divided into training and tested sections using a variety of machine learning methods. Early detection of renal issues lowers the risk of CKD. Algorithms are used to forecast kidney issues. The k-nearest neighbor, decision tree classifier, support vector classifier, logistic regression, random forest classifier, and extra tree classifier are some of the classification methods used on the Jupiter platform. The best and most effective method for detecting renal disease is this one. The random forest classifier is the best binary classifier for predicting renal illness when it comes to diagnosis[8][9].

MATERIAL AND METHODS

This study's main impartial is to investigate the accuracy of CKD detection using feature selection and machine learning methodologies. The identified classification method can provide predicted values for an early CKD diagnosis.





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The Proposed Framework

The Recommended Framework Figure 1.1 shows the suggested methodology for creating identification machine learning models and a comparison of them. The main objective of this learning is to provide a machine erudition approach using multiple classification algorithms for CKD identification. To diagnose CKD early, it can be helpful to find approaches with a high percentage of successfully classified examples and acknowledged classifiers. In this comparison analysis, the suggested method is compared to various state-of-the-art approaches.

Source of the Data & Description

The dataset for this study is based on patient data for chronic renal disease obtained from the Kidney Care Centre inNila Nursing HomeTamil Nadu, India. In addition to 13 numerical and 14 nominals, the dataset contains 1100 observations of individuals with the condition. The dataset, which quantifies the severity of the renal disease, consists of 275 observations at stage 1, 269 observations at stage 2, 133 observations at stage 3A, 109 observations at stage 3B, 145 observations at stage 4, and 169 observations at stage 5.Among the data variables in the dataset are those about Id, Age, Gender, Blood pressure, Anaemia, Blood Glucose Random, Specific gravity, Pus Cell, Albumin, Sugar, Diabetes mellitus, Packed Cell Volume, Sodium, Appetite, Hypertension, Bacteria, Red Blood Cells, Potassium, Pedal Edema, Red blood cell count, Serum creatinine, Blood Urea, Haemoglobin, Pus Cell Clumps, White blood cell count, Coronary Artery Disease, and CKD Stages.

Data preprocessing

The dataset is chosen for CKD prediction to facilitate data analysis and useful knowledge. To deploy a machine learning classifier for a chosen dataset, sufficient data must be available. The dataset has 27 attributes and is organized in attribute-relation file format. To use associative approaches, the dataset is transformed into a binomial format. Additionally, there are duplicate records and missing records. Category variables are converted to numeric values via label encoding.

Feature Selection

To reduce overfitting and increase projected accuracy, this class provides a meta-estimator that fits multiple randomized decision trees (extra-trees) to different dataset subsamples. The data set was split into two parts before modelling could start: The Train set (70%) was used to select and validate models, and the Test set (30%) contained data used to determine how effectively models generalized to new data. Data for instruction: The training dataset, which is derived from the main dataset, has 770 of the 1100 elements in the CKD main dataset. Information for testing: Out of the 1100 entries in the core CKD dataset, 330 were used in the testing.

Machine learning Model

This work focuses on developing a cataloguing model with feature selection based on evidence obtained to classify chronic kidney disease (CKD) using the Ensemble Random Forest classification technique. To function as an ensemble classification process, the random forest process builds a variety of decision trees. Random forests' hyper parameters bear striking similarities to decision trees' and bagging classifiers' hyper parameters. It is not necessary to combine a decision tree with a bagging classifier when using the random forest's classifier class. Using a random forest and the algorithm's regressor, regression jobs may be managed. While the trees are growing, the random forest improves the model's arbitrariness. Instead of looking for the main component when dividing a hub, it looks for the best part from an erratic subset of supply. As a result, there are many possibilities, the majority of which provide a better model. The training data sets' random subsets are used to build several decision trees. A vast collection of decision trees offers better outcome accuracy. The algorithm's runtime is reasonably quick and it takes into account missing data. The method is randomly generated by Random Forest, not the training set. The decision class is a type of class that decision trees produce.





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Entropy

Entropy is a machine learning metric that quantifies the level of disorder or uncertainty in a system or dataset. It is a statistic that measures the amount of data in a dataset and is frequently used to evaluate the accuracy and quality of a model's predictions.

```
Entropy = -p(a) * \log(p(a)) - p(b) * \log(p(b))
```

Throughout the dataset, we have observations from Stages 1 through 5, including Stages 1, 2, Stages 3A and 3B. Then, according to the following equation: $E = -(P_j \log_2 p_j + P_k \log_2 p_k + P_l \log_2 p_l + P_m \log_2 p_m + P_n \log_2 p_n + P_l \log_2 p_n)$

Where P_i is the probability of selecting patients in Stage 1;

 P_k denotes the probability of selecting patients in Stage 2;

 P_l denotes the likelihood of selecting patients in Stage 3A.

 P_m = Probability of selecting Stage 3B patients;

 P_n = Probability of selecting Stage 4 patients and;

 P_0 = Probability of selecting Stage 5 patients.

Information Gain

After dividing a dataset on an attribute, the reduction in entropy is used to determine the information gain. The goal of building a decision tree is to locate the characteristic that provides the largest information gain (i.e., the most homogenous branches). The pattern detected in the dataset and the reduction in entropy is termed information gain. The following formula may be used to describe knowledge gain mathematically:

```
Information\ gain = E(parent) - [weighted\ average]E(children)
```

Each of our n datasets has m entries. The following pseudocode, which looks like Python, illustrates bootstrap sampling:

Pseudocode of an ensemble for multi-class classification using random forest:

```
mc_models = []
    for i in range (n_estimators):
    erfmodel = erf (max_depth=10, random state = i)
    mc_models.append(erfmodel)

ckdstage_prediction = []
    for erfmodel in mc_models:
    y_pred = mc_models.predict(x_test)
    ckdstage_prediction.append (y_pred)
```

Prediction of the Stages of CKD

We establish the important measures to evaluate the model's performance. We may develop measurement metrics to represent the performance of the classifiers in the digital diagnostics challenge using four essential characteristics.

Multi-Classification Confusion Matrix for Algorithms

```
r = assigned label

c = prediction classification

for each instance of the input do

K(r,c) += 1

end for
```

K is the confusion matrix, where *c* and *r* stand for the matrix's column and row, respectively.





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Accuracy of Classification

The created classifier model's accuracy may be designed using the equation below.

$$Accuracy = \frac{TN + TP}{TP + FP + TN + FN}$$

Where TP = Predicted is likewise positive and observation is positive

TN: Both the observed and the projected outcomes are negative.

FP = Predicted is positive while observation is negative

FN stands for "Observation is positive, but prediction is negative".

RESULTS AND DISCUSSION

After successfully the dataset through multi-classification machine learning algorithms,we acquired the accuracy numbers mentioned below. All metrics true positive, precision, or any other metric are calculated in multi-class classification in the same way they are in binary, with the exception that they must be calculated for each class. If we compute the true positive, false positive, true negative, and false negative given a class, we may virtually derive any metric for that class. In multi-class problems, precision has been calculated for each class, whereas in binary class problems, we only have one number. One global measure may be used to evaluate multi-class data and get micro, macro, and weighted precision. By adding micro, macro, and weighting, any metric from the confusion matrix may be transformed into a global metric. The figure 4.1 classifiers successfully predicted the CKD Stages at Stage 1 scoring 71 out of 74, and 4 of those indicated a poor result. The CKD Stage 2 is 66 out of 67, with 1 bad consequence anticipated. The CKD Stage 3A is 40 out of 42, and 2 outcomes were projected to be unfavourable. The CKD Stage 3B is 31 out of 33, and 2 outcomes were projected to be unfavourable. The CKD Stage 4 is 46 out of 51 and 5 outcomes were projected to be unfavorable. The CKD Stage 5 is 63 out of 63, meaning that not a single aspect in the report accurately indicated a bad result. The figure 4.2 classifiers successfully predicted the CKD Stages at Stage 1 scoring 72 out of 74, and Stage 2 was expected to be unsuccessful.

The CKD Stage 2 score is 67 out of 67, with 0 unfavourable outcomes foreseen. The CKD Stage 3A is 39 out of 42, and 3 outcomes were projected to be unfavourable. The CKD Stage 3B is 31 out of 33, and 2 outcomes were projected to be unfavourable. The CKD Stage 4 is 46 out of 51 and 5 outcomes were projected to be unfavorable. The CKD Stage 5 is 63 out of 63, meaning that not a single aspect in the report accurately indicated a bad result. The figure 4.3 classifiers successfully predicted the CKD Stages at Stage 1 scoring 72 out of 74, and Stage 2 was expected to be unsuccessful. The CKD Stage 2 is 66 out of 67, with 1 bad consequence anticipated. The CKD Stage 3A is 41 out of 42, and 1 unfavorable result was expected. The CKD Stage3B is 31 out of 33, and 2 outcomes were projected to be unfavourable. The CKD Stage 4 is 46 out of 51 and 5 outcomes were projected to be unfavorable. The CKD Stage 5 is 63 out of 63, meaning that not a single aspect in the report accurately indicated a bad result. The figure 4.4 classifiers correctly predicted the CKD Stages, with Stage 1 scoring 72 out of 74 and Stage 2 predicting a poor result. The CKD Stage 2 score is 67 out of 67, with 0 unfavourable outcomes foreseen. The CKD Stage 3A is 41 out of 42, and 1 unfavorable result was expected. The CKD Stage 3B is 31 out of 33, and 2 outcomes were projected to be unfavourable. The classifiers correctly predicted the CKD Stages, with Stage 4 being 46 out of 51 and 5 predicting a bad result. The classifiers successfully predicted the CKD Stages at Stage 5 had a score of 63 out of 63, and 0 of those predictions were incorrect across the board. The results of the validation accuracy for the multi-class classifier outlined above are summarized in Figure 4.5 overall in the CKD stage.

CONCLUSION

One of the difficult research issues in medical science nowadays seems to be the accurate prognosis of chronic renal disease. To identify which patients will exhibit which CKD symptoms at an early stage, this research proposes a prediction method. The models are trained and verified using the input parameters provided in the dataset, which include input parameters gathered from CKD patients. To perform the CKD diagnosis, multi-class classification with





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ensemble random forest machine learning models is created. Out of the 26 CKD stages, the overall CKD stage accuracy for the 26 characteristics is 96.06%. Out of the 26 CKD stages, the overall CKD stage accuracy for the 20 characteristics is 96.36%. Out of the 26 CKD Stages, the accuracy of the CKD Stages for 15 features is 96.66% overall. Finally, for the multiclass classification out of the 26 CKD stages, the overall CKD stage accuracy for the 10 features is 97%.

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Table 1: Multi-Class Classification using Ensemble Random Forest to classify datawith 26 features, 20 features, 15 features, and 10 features Classification Report

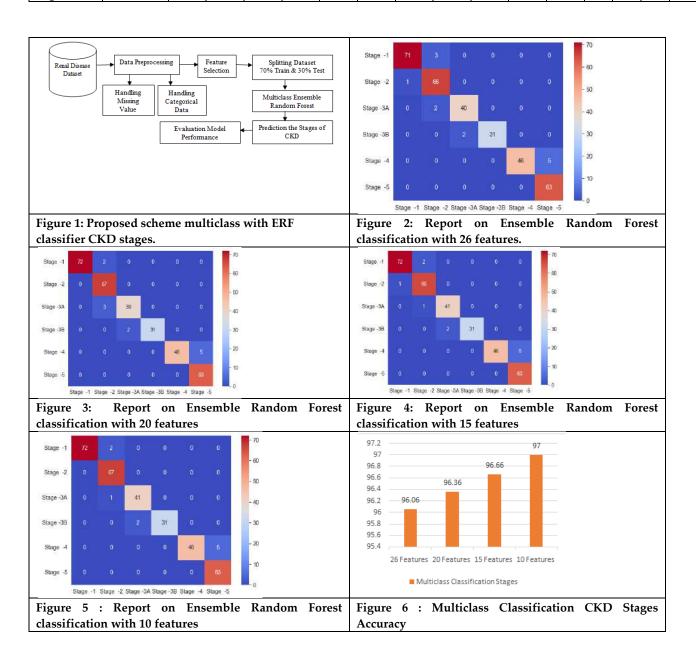
	Precision				Recall			F1-Score				Support				
Features	26	20	15	10	26	20	15	10	26	20	15	10	26	20	15	10
Stage 1	0.99	1.00	0.99	1.00	0.96	0.97	0.97	0.97	0.97	0.99	0.98	0.99	74	74	74	74
Stage 2	0.93	0.93	0.96	0.96	0.99	0.93	0.99	1.00	0.96	0.96	0.97	0.98	67	67	67	67
Stage 3A	0.95	0.95	0.95	0.95	0.95	0.95	0.98	0.98	0.95	0.94	0.96	0.96	42	42	42	42
Stage 3B	1.00	1.00	1.00	1.00	0.94	1.00	0.94	0.94	0.97	0.97	0.97	0.97	33	33	33	33
Stage 4	1.00	1.00	1.00	1.00	0.90	1.00	0.90	0.90	0.95	0.95	0.95	0.95	51	51	51	51
Stage 5	0.93	0.93	0.93	0.93	1.00	0.93	1.00	1.00	0.96	0.96	0.96	0.96	63	63	63	63
Accuracy	0.96	0.96	0.97	0.97		330	330	330	330							
Macro avg	0.97	0.97	0.97	0.97	0.96	0.96	0.96	0.97	0.96	0.96	0.97	0.97	33	33	33	33





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													0	0	0	0
Weighted													33	33	33	33
avg	0.96	0.97	0.97	0.97	0.96	0.96	0.97	0.97	0.96	0.96	0.97	0.97	0	0	0	0







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RESEARCH ARTICLE

Eccentricity Energy of Cover Pebbling Graphs

C. Muthulakshmi@Sasikala¹ and A. Arul Steffi²

¹Associate Professor of Mathematics, Sri Paramakalyani College, Alwarkurichi, Tenkasi, India.

²Research Scholar of Mathematics, St. Xavier's College (Autonomous), Palayamkottai, Tirunelveli, India.

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*Address for Correspondence

C. Muthulakshmi@Sasikala Associate Professor of Mathematics, Sri Paramakalyani College, Alwarkurichi, Tenkasi, India. Email: kalasasispkc@gmail.com



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ABSTRACT

Eccentricity matrix $E_{co}(G)$ of cover pebbling graph G is defined from the weight of an edge e = uvbetween two vertices u and v. Eccentricity eigen values of a graph G are the eigen value of its eccentricity matrix $E_{cp}(G)$. The eccentricity spectrum of a cover pebbling graph G consists of the eccentricity eigen values of its eccentricity matrix $E_{cp}(G)$. Eccentricity energy $E_{cp}(E)$ is summation of absolute values of eccentricity eigen values of the eccentricity matrix $E_{cp}(G)$. In this paper, we study the eccentricity energy $E_{cp}(E)$ of cover pebbling complete graph, cover pebbling path graph, cover pebbling bipartite graph.

Keywords: Eccentricity matrix, Eccentricity eigen values, Eccentricity energy, cover pebbling graph.

INTRODUCTION

Let G = (V(G), E(G)) be any simple and connected graph with n vertices and m edges. If $u, v \in V(G)$ and $uv \in E(G)$, a pebbling move is the removal of two pebbles from vertex u and placing one pebble on an adjacent vertex v throwing one pebble away. Define the weight of an edgee, w(e) as follows.

w(u, v) = 1 if $uv \in E(G)$ and a pebbling move occurs between u and v.For a vertex v of G, the pebbling eccentricity of a vertex v is $e_p(v) = \max\{w(u, v); u \in V(G)\}$. The eccentricity matrix $E_{cp}(G) = (a_{ij})$ is defined as follows;

$$E_{cp}(G) = (a_{ij}) = \begin{cases} w(v_i, v_j) & \text{if } w(v_i, v_j) = \min\{e_p(v_i), e_p(v_j)\} \\ 0 & \text{otherwise} \end{cases}$$

Eccentricity matrix of graphs are discussed by Wang in [7]. Graph pebbling was first introduced by Lagarias and Saks and Chung [2] established various results concerning pebbling numbers. Sakunthala Srinivasan and Vimala Shanmugavel[5] introduced the concept of pebbling graph. Cover pebbling graphs were introduced in [4]. Energy of a graph is from Gutman[3]. In this paper, we combine the concepts of cover pebbling graphs and eccentricity matrix.





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Preliminaries

Definition 2.1. In a simple graph, consisting of two vertices, a pebbling move [4] can be defined as the removal of two pebbles from the first vertex and placing one pebble on an adjacent vertex.

Definition 2.2. Define the weight of an edge e = uv between two vertices u and v as follows: w(e) = w(uv) = v(v)Iif $uv \in E(G)$ and a pebbling move occurs between u and v. The number of pebbling move occurs between u and v is the weight of the edge connecting the vertices. Let $P_{ij} = v_i, v_{i+1}, v_{i+2}, \dots, v_{j-1}, v_j$ be the shortest $v_i - v_j$ path. Then $w(P_{ij}) = \sum_{l=i}^{j-1} w(v_l, v_{l+1}).$

Definition 2.3. In a simple graph, consisting of two vertices, place a pebble on every vertex of the graph using sequence of pebbling moves is called a cover pebbling graph[4].

Definition 2.4. Energy graph E(G) is the total of absolute values of the eigen values $\lambda_1, \lambda_2, \dots, \lambda_n$ of the adjacency matrix and it is expressed as $E(G) = \sum_{i=1}^{n} |\lambda_i|$.

Definition 2.5. The eccentricity energy of the cover pebbling graph G is defined by $E_{cp}(E) = \sum_{i=1}^{n} |\lambda_i|$ where $\lambda_1, \lambda_2, \cdots, \lambda_n$ are the eccentricity eigen values of eccentricity matrix of the corresponding cover pebbling graph G.

Definition 2.6. A complete graph is a simple graph in which any two vertices are adjacent and a complete graph with n vertices is denoted by K_n.

Definition 2.7. A path on n-vertices denoted by P_n is a sequence of vertices of which any two are adjacent if they are consecutive, and are nonadjacent otherwise.

Definition 2.8. A graph is bipartite if its vertices can be partitioned into two subsets X and Y so that every edge has one end in X and other end in Y. If |X| = n and |Y| = n, then complete bipartite graph is denoted by $K_{n,n}$.

Theorem 2.9. The cover pebbling number of complete graph K_n is 2n - 1.[1]

Theorem 2.10. The cover pebbling number of path P_n is $2^n - 1$.[1]

Theorem 2.11. The cover pebbling number of complete bipartite graph $K_{n,n}$ is 6n - 3.[6]

MAIN RESULTS

Example 3.1. Eccentricity energy of cover pebbling complete graph K_4 . Now, $w(v_1, v_2) = 1$, $w(v_2, v_1) = 1$, $w(v_3, v_1) = 1$ $1, w(v_4, v_1) = 1, w(v_1, v_3) = 1, w(v_2, v_3) = 0, w(v_3, v_2) = 0, w(v_4, v_2) = 0, w(v_1, v_4) = 1, w(v_2, v_4) = 0, w(v_3, v_4) = 0, w(v_3, v_4) = 0, w(v_4, v_4) = 0, w$ $0, w(v_4, v_3) = 0, e_p(v_1) = 1, e_p(v_2) = 1, e_p(v_3) = 1, e_p(v_4) = 1.$

 $\min\{e_p(v_1), e_p(v_2)\} = 1 = w(v_1, v_2)$

Hence $a_{12} = 1$.

$$\min\{e_p(v_1),e_p(v_3)\}=1=w(v_1,v_3).$$

Hence $a_{13} = 1$ and so on.

Also
$$a_{11} = a_{22} = a_{33} = a_{44} = 0$$
.

Also $a_{11} = a_{22} = a_{33} - a_{44} - 3$. Eccentricity matrix $E_{cp}(G) = \begin{pmatrix} 0 & 1 & 1 & 1 \\ 1 & 0 & 0 & 0 \\ 1 & 0 & 0 & 0 \\ 1 & 0 & 0 & 0 \end{pmatrix}_{4\times4}$

Characteristic equation is $det(E_{cp}(G) - \lambda I) = 0i.e., \lambda^2(\lambda^2 - 3) = 0.$

 $\lambda = 0$ (two times) and $\lambda^2 = 3$.





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 $\lambda = \pm \sqrt{3}$

Eccentricity energy = $2\sqrt{3}$.

Theorem 3.2. Let K_n be the complete graph with n vertices and m edges. Then eccentricity energy of cover pebbling complete graph K_n is $2\sqrt{n-1}$ where n > 2.

Proof. Let K_n be the complete graph with n vertices $\{v_1, v_2, \dots, v_n\}$ and m edges. Weight of an edge between v_1 to any other vertex is one. That is $w(v_1, v_2) = w(v_2, v_1) = 1$. In general, $w(v_1, v_1) = w(v_1, v_1) = 1$ where $i = 2, 3, \dots, n$. But weight of any edge between any two vertices v_i and v_j where $2 \le i, j \le n$ is zero. Now, $\min\{e_p(v_1), e_p(v_j)\} = w(v_1, v_j) = 1 \text{ where } j = 2, 3, \dots, n$ $\min\{e_p(v_i), e_p(v_i)\} \neq w(v_i, v_i) \text{ where } 2 \leq i, j \leq n.$

Eccentricity matrix for cover pebbling complete graph G is

$$E_{cp}(G) = \begin{pmatrix} 0 & 1 & & \cdots & 1 \\ 1 & 0 & 0 & \cdots & 0 \\ 1 & 0 & 0 & \cdots & 0 \\ \vdots & \vdots & \vdots & \ddots & \vdots \\ 1 & 0 & 0 & \cdots & 0 \end{pmatrix}_{n \times n}.$$

The characteristic equation is $\lambda^{n-2} (\lambda^2 - (n-1)) = 0$.

The eccentricity eigen values are 0 (n-2) times and $\lambda = \pm \sqrt{n-1}$. Eccentricity energy $E_{cp}(E) = 2\sqrt{n-1}$.

Example 3.3. Eccentricity energy of cover pebbling path graph P₄. Now,

$$\begin{array}{l} w(v_1,v_2) = 7, w(v_2,v_1) = 7, w(v_3,v_1) = 10, w(v_4,v_1) = 11, w(v_1,v_3) = 10, \\ w(v_2,v_3) = 3, w(v_3,v_2) = 3, w(v_4,v_2) = 4, w(v_1,v_4) = 11, w(v_2,v_4) = 4, \\ w(v_3,v_4) = 1, w(v_4,v_3) = 1, e_p(v_1) = 11, e_p(v_2) = 7, e_p(v_3) = 10, e_p(v_4) = 11. \\ \min\{e_p(v_1), e_p(v_2)\} = \min\{11,7\} = 7 = w(v_1,v_2). \end{array}$$

Hence $a_{12} = 7 = a_{21}$.

$$\min\{e_n(v_1), e_n(v_3)\} = \min(11,10) = 10 = w(v_1, v_3).$$

Hence $a_{13} = 10 = a_{31}$

$$\min\{e_p(v_1), e_p(v_4)\} = 11 = w(v_1, v_4).$$

Hence $a_{14} = 11 = a_{41}$. Other $a_{ij} = 0$ where $2 \le i, j, \le 4$.

Again $a_{11} = a_{22} = a_{33} = a_{44} = 0$.

Eccentricity matrix
$$E_{cp}(G) = \begin{pmatrix} 0 & 7 & 10 & 11 \\ 7 & 0 & 0 & 0 \\ 10 & 0 & 0 & 0 \\ 11 & 0 & 0 & 0 \end{pmatrix}_{4\times4}$$

Characteristic equation is $det(E_{cn}(G) - \lambda I) = 0$

i. e.,
$$\lambda^2(\lambda^2 - 270) = 0$$
.

 $\lambda = 0$ (two times) and $\lambda = \pm \sqrt{270}$.

Eccentricity energy = $2\sqrt{270}$.

$$\begin{array}{l} \text{ Theorem 3.4. Let } P_n \text{ be a path on } n \text{ vertices } (n \geq 3). \text{ Then eccentricity energy of } P_n \text{ is } \\ E_{cp}(E) = 2\sqrt{\sum_{j=1}^{n-2}\Bigl(\sum_{i=1}^{j}(2^{n-i}-1)\Bigr)^2 + [\sum_{k=1}^{n-2}(2^{n-k}-1)+1]^2}. \end{array}$$

Proof. Let P_n be a path on n vertices $\{v_1, v_2, \cdots, v_n\}$ where $n \ge 3$.

$$\begin{array}{l} w(v_1,v_2) = w(v_2,v_1) = 2^{n-1} - 1 \\ w(v_1,v_3) = w(v_3,v_1) = 2^{n-1} - 1 + 2^{n-2} - 1 \\ \vdots \\ w(v_1,v_n) = w(v_n,v_1) = 2^{n-1} - 1 + 2^{n-2} - 1 + \dots + 2^{n-(n-2)} - 1 + 1 \\ e_p(v_1) = 2^{n-1} - 1 + 2^{n-2} - 1 + \dots + 2^{n-(n-2)} - 1 + 1 \\ w(v_2,v_3) = w(v_3,v_2) = 2^{n-2} - 1 \end{array}$$





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$$\min(e_p(v_i), e_p(v_j)) \neq w(v_i, v_j)$$

Hence $a_{ij} = 0 = a_{ji}$.

Also, $a_{ii} = 0$ where $i = 1, 2, 3, \dots, n$.

Eccentricity matrix of cover pebbling path P_n is

$$E_{cp}(P_n) = \begin{pmatrix} 0 & 2^{n-1} - 12^{n-1} - 1 + 2^{n-2} - 1 \cdots a & b \\ 2^{n-1} - 1 & 0 & 0 & \cdots 0 & 0 \\ 2^{n-1} - 1 + 2^{n-2} - 1 & 0 & 0 & \cdots 0 & 0 \\ \vdots & \vdots & \vdots & \ddots & \vdots & \vdots \\ a & 0 & 0 & \cdots 0 & 0 \\ b & 0 & 0 & \cdots 0 & 0 \end{pmatrix}$$
and b= $\sum_{n=1}^{n-1} (2^{n-i} - 1)$

where $a = \sum_{i=1}^{n-2} (2^{n-i} - 1)$ and $b = \sum_{i=1}^{n-1} (2^{n-i} - 1)$.

The characteristic equation is

$$\lambda^{n-2}(\lambda^2 - \left[\sum_{j=1}^{n-2} \left(\sum_{i=1}^{j} (2^{n-i} - 1)\right)^2 + \left[\sum_{k=1}^{n-2} (2^{n-k} - 1) + 1\right]^2\right] = 0$$

 $\lambda = 0 (n-2)$ times and

$$\begin{split} \lambda^2 &= \sum_{j=1}^{n-2} \Biggl(\sum_{i=1}^j \bigl(2^{n-i} - 1 \bigr) \Biggr)^2 + \Biggl[\sum_{k=1}^{n-2} \bigl(2^{n-k} - 1 \bigr) + 1 \Biggr]^2 \\ \lambda &= \pm \sqrt{\sum_{j=1}^{n-2} \Biggl(\sum_{i=1}^j \bigl(2^{n-i} - 1 \bigr) \Biggr)^2 + \Biggl[\sum_{k=1}^{n-2} \bigl(2^{n-k} - 1 \bigr) + 1 \Biggr]^2} \end{split}$$

 $\text{Eccentricity energy,} E_{cp}(E) = 2\sqrt{\sum_{j=1}^{n-2} \Bigl(\sum_{i=1}^{j} (2^{n-i}-1)\Bigr)^2 + \bigl(\sum_{k=1}^{n-2} (2^{n-k}-1) + 1\bigr)^2}.$

Example 3.5. Eccentricity energy of cover pebbling complete bipartite graph $G = K_{2,2}$.

Eccentricity matrix
$$E_{cp}(G) = \begin{pmatrix} 0 & 3 & 4 & 1 \\ 3 & 0 & 0 & 0 \\ 4 & 0 & 0 & 0 \\ 1 & 0 & 0 & 0 \end{pmatrix}_{4 \times 4}$$

Characteristic equation is $\lambda^2(\lambda^2 - 26) = 0$.

$$\lambda = 0$$
 (2 times), $\lambda = \pm \sqrt{26}$.

Eccentricity energy, $E_{cp}(E) = 2\sqrt{26}$.

Theorem 3.6. Let $K_{n,n}$ be the complete bipartite graph with 2n vertices where $n \ge 2$. The eccentricity energy of cover pebbling complete bipartite graph G is $E_{cp}(E) = 2\sqrt{n(4n^2 - 3)}$ where $n \ge 2$.





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 $\textbf{Proof}. \text{ Let } K_{n,n} \text{ be the complete bipartite graph with vertex sets } X = \{u_1, u_2, \cdots, u_n\} \text{ and } Y = \{v_1, v_2, \cdots, v_n\}. \text{ Eccentricity are sets } X = \{u_1, u_2, \cdots, u_n\} \text{ and } Y = \{v_1, v_2, \cdots, v_n\}. \text{ Eccentricity }$ matrix of cover pebbling complete bipartite graph G is

$$E_{cp}(G) = \begin{pmatrix} 0 & 2n-1 & 2n & 1 & 2n & \cdots & 1 \\ 2n-1 & 0 & 0 & 0 & 0 & 0 & \cdots & 0 \\ 2n & 0 & 0 & 0 & 0 & 0 & \cdots & 0 \\ 1 & 0 & 0 & 0 & 0 & 0 & \cdots & 0 \\ 2n & 0 & 0 & 0 & 0 & 0 & \cdots & 0 \\ 2n & 0 & 0 & 0 & 0 & 0 & \cdots & 0 \\ 1 & 0 & 0 & 0 & 0 & 0 & \cdots & 0 \\ 2n & 0 & 0 & 0 & 0 & 0 & \cdots & 0 \\ 2n & 0 & 0 & 0 & 0 & 0 & \cdots & 0 \\ \vdots & \vdots & \vdots & \vdots & \vdots & \vdots & \vdots & \vdots & \vdots \\ 1 & 0 & 0 & 0 & 0 & 0 & 0 & \cdots & 0 \end{pmatrix}$$
 The characteristic equation is $\lambda^{2n-2} \left(\lambda^2 - \left((2n-1)^2 + (n-1)(2n)^2 + (n-1) \right) \right) = 0$.

Eccentricity eigen values are $\lambda = 0(2n-2)$ times and $\lambda^2 = (2n-1)^2 + (n-1)(2n)^2 + (n-1)$.

$$\lambda = \pm \sqrt{n(4n^2 - 3)}.$$

Eccentricity energy , $E_{cp}(E) = 2\sqrt{n(4n^2 - 3)}$ where $n \ge 2$.

CONCLUSION

The eccentricity energy of cover pebbling complete graph, cover pebbling path and cover pebbling bipartite graph have been discussed. The eccentricity energy of cover pebbling cycle is an open problem.

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RESEARCH ARTICLE

Implementation of Time series Control charts using PCA and ARIMA Model

R. Sasikumar* and M.Sujatha

Department of Statistics, Manonmaniam Sundaranar University, Tirunelveli, Tamil Nadu, India.

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*Address for Correspondence

R. Sasikumar

Department of Statistics, Manonmaniam Sundaranar University, Tirunelveli, Tamil Nadu, India. Email: sasikumarmsu@gmail.com



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ABSTRACT

Public healthcare aims to promote and protect the health factors and environmental factors that contribute to the safety and overall health of persons and communities. It involves the analysis and management of data related to various health determinants, such as air; water, etc...environmental exposures, and healthcare services utilization. SPC (Statistical process control chart) is the procedure for monitoring the process and control with the help of statistical tools and techniques. ARIMA (Autoregressive integrated moving averages) control chart is a type of SPC time series chart that can be used to observe and handle processes over time. To determine the process signals out of control and trends or patterns that might be present in the data, process control charts are used, particularly ARIMA control charts are engaged. The secondary data contains various air pollutant factors like PM 2.5, PM 10, nitrogen dioxide(NO2), sulfur dioxide(SO2), carbon monoxide (CO) and other relevant pollutants from major cities in Tamil Nadu. This research contains three parts (i) Tirupur city identified as the most pollutant city (ii) the PCA (principal component analysis) technique used to determine the significant components (iii) estimating the parameters with ARIMA model. The results showed that the SCC (Special cause control chart) is more appropriate for auto-correlated data to evaluate the stability of the data since it provides a higher probability of coverage than individual control charts.

Keywords: ARIMA control charts, Public health, control charts, PCA





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INTRODUCTION

Public health is the term used for protecting, upgrading, hopeful and administration of healthcare factors and environmental factors. To promote and protect community health, policymakers, researchers, and public health workers can use information about public health to make well-informed decisions, create evidence-based plans, and carry out actions. Air quality is one of the major determinants of public health. Most cities worldwide experience severe air-quality problems, and they have received rising attention in the past decade (Vallero.D.A, 2014). Toxic air pollutants have two types: Physical and Chemical properties. Specified pollutants are those that are common in the United States and are known to be harmful to public health and the environment, as defined by the Clean Air Act of 1971. There are six major pollutants: lead, ozone, sulphur dioxide, nitrogen dioxide and Particular matters sizeunder 10 and 2.5 (Mangayarkarasi, R, et al. 2021). Particulate matter (PM): Airborne particles like small dust, filth and smoke are known as PM 10 and PM 2.5. When inhaled these particles, especially the tiny ones known as PM 2.5 and PM 10 can vary in size and composition and harm health. Nitrogen oxide (NO_x) is gases that are essentially created during the combustion process such as when fossil fuels are used in automobiles, power plants and industrial settings. They can have negative effects on the cardiovascular and respiratory systems. Sulphur dioxide (SO₂): The burning of fossil fuels emits gas that contains sulfur, such as coal and oil. It is a major provider of acid rain. Carbon Monoxide (CO): The incomplete combustion of fossil fuels, largely in vehicles and industrial processes produced this colourless and odourless gas. Ozone (O3): is a gas that protects humans from damaging UV rays. Lead (Pb): Lead is a poisonous heavy metal that was traditionally used in gasoline, paints and other products (Kokilavani.S, et al. 2020). A quantitative metric of all the pollutants is known as the Air quality index (AQI). Higher AQI values denote a hazardous level of exposure for the general public (Sasikumar.R, et al. 2022).

SPC is a methodology for process monitoring and control that makes use of statistical tools and techniques. One of the most important problem-solving instruments for monitoring stability in processes and enhancing capability through decreased variability is the control chart (Montgomery D.C. 2009). This paper includes data from all the major cities of Tamil Nadu. Tirupur is identified as a major pollutant city. Due to a variety of anthropogenic activities and natural reasons, this city experiences air pollution problems. Autocorrelation in observations makes it difficult to identify "special causes". There is a large chance of false positives or negatives due to autocorrelation in observations, which makes it challenging to distinguish between "special causes" that are present and those that are not. (Smeti et al.2007a). After selecting and adjusting an ARIMA model to the process, the residuals are examined using a conventional control chart. It makes reason to observe residuals instead of actual observations because residuals are uniformly distributed with an average of zero if the process is within control and stays independent of possible mean differences when the process differs from control (Russo, et al 2012). Since several quality characteristics can be affected by the air, for this study we choose only two quality characteristics by using the PCA technique. Principal components are used to determine which variables or groups of variables have the greatest influence to detect a likelihood of an out-of-control signal in order to subsequently identify it as a primary operating source(Souza, A. M. et al 2012) This research paper aims to compare the traditional control charts and residuals from the fitted ARIMA control charts.

METHODOLOGY

The methods included in this study are

- Principal Component Analysis(PCA).
- Statistical Process Control charts.

Principal Component Analysis (PCA)

PCA is a dimensionality reduction technique that enables us to reduce the key information from a big collection of variables into a more approachable collection of main components, which are uncorrelated variables. We successfully minimize the dimensionality of the dataset while maintaining the most important data, by choosing the





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principal components with the highest eigen values or the greatest proportion of variance explained (Karamizadeh.S et al 2013)

The steps that involve

• Standardization is the initial step, ensuring that each feature in the dataset has a mean (average) of 0 and a standard deviation of 1.The formula for standardization

 $Z=X-\mu/\sigma$ (1)

where Z represents the standard value

X represents the values

μ represents the mean

σ represents the standard deviation

- Calculate the covariance matrix (C); Eigen values (L) and eigenvectors (λ L).
- Select a subset of the top eigenvectors and their corresponding eigenvalues based on the explained variance. These eigenvectors become the principal components.

Control charts

Traditional control charts frequently make the assumption that observations pertaining to processes have an independent distribution with a constant mean and variance. Data autocorrelation makes it more difficult to distinguish between "special causes" that exist and those that do not, increasing the risk of incorrect results and/or false positives. (Montgomery 2007).

Autocorrelation (ACF) and Partial autocorrelation function (PACF)

The term "autocorrelation" refers to a measure of interdependence between the observations. Data with a self-correlation are automatically correlated. The primary result of autocorrelation in data analysis on control charts is the production of far tighter than ideal control limits. This means that the prior approach needs to be modified to allow for the possibility of sufficient autocorrelation in the process data (Elevli.S, et al 2016). Data autocorrelation can be detected using the ACF and PACF. The association between data points in a time series and the previous series is plotted using the ACF method. The relationship between the time series variables at time t and all other values at that time (t-1) is represented by the first lag. Once interval delays are taken into account, PACF computes the correlation between data points at time t and those at t-1 (Russo, et al 2012).

Special cause control charts(SCC)

Establishing accurate control limits in control charts is one of the essential steps to ensure the statistical consistency of the process under analysis. If there are different levels of autocorrelation in the quality feature under investigation, the control charts derived using the independence assumption will display more false alarms. When data from control charts are evaluated, autocorrelation mostly results in considerably tighter control limits than planned. If the process data exhibits a significant amount of autocorrelation, the traditional approaches must be changed to consider this. In order to create control charts for auto-correlated data, there are two main methods. The first method uses traditional control charts but modifies the technique for calculating the process variance to estimate the actual process variation. As a result, the correlation in this chart is handled by positioning the control boundaries by the altered variability. The second effective approach for handling auto-correlated data is to employ a suitable time series model to explicitly characterize the correlative structure. Then, using that model, remove autocorrelation from the data and apply control charts to the residuals (Montgomery 2007). An SCC chart is the name for this form of control chart, which plots the residuals of the ARIMA model after fitting it to the process. This approach of graphics is compatible with all conventional process control instruments. The centre line is 0 because the mean of the residuals is nil. The residual's standard deviation σ_a is the standard deviation employed in this situation. The chart limits are determined by (i) the upper control limit of 3.0a (ii) the centre line of 0 and (iii) the lower control limit of 3.σ_a.(Perzy.M,et al 2015).





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Auto Regressive Integrated Moving Average(ARIMA)

The famous time series forecasting technique known as ARIMA is used to analyze and predict data that has temporal relationships. It combines moving average (MA), autoregressive (AR), and differencing (I) components. The use of the variable's historical values to forecast its future values is referred to as the AR component. It is predicated on the idea that a variable's present value is linearly related to its past values. The time series is made stationary by the employment of the I component, which stands for differencing. The stationarity assumption, which holds that the statistical properties of the data do not change over time, is an essential basis of time series analysis. Differencing is taking the current value and subtracting it from the previous value in order to get rid of seasonality and patterns. The MA element is for error terms of residuals (**Kovarik.M et al 2015**). The ARIMA model's order is determined by three variables: p, d, and q. the parameters p and q stand for the moving average components and the autoregressive components lagged error terms, respectively, while d stands for the order of differencing needed to prove stationarity.

 $Y\left(t\right) = C + \Phi_{1} * Y_{(t-1)} + \Phi_{2} * Y_{(t-2)} + ... + \Phi_{p} * Y_{(t-p)} + \theta_{1} * \varepsilon_{(t-1)} + \theta_{2} * \varepsilon_{(t-2)} + ... + \theta_{p} * \varepsilon_{(t-q)} + \varepsilon_{(t)}$

where, the time series at time t is represented by Y(t). The expression C is constant.

- The autoregressive coefficients Φ_1 , Φ_2 ... Φ_p indicate how dependent the time series is on earlier values.
- $Y_{(t-1)}$, $Y_{(t-2)}$... and $Y_{(t-p)}$ represent the time series' lagged values.
- The moving average coefficients denoting the reliance on previous error terms are represented by the values of θ_1 , θ_2 ... θ_p .
- The lagged error terms are ε (t-1), ε (t-2)... ε (t-q).

The current error term, denoted as ϵ (t), is presumed to have a normal distribution with a mean of zero and a constant variance.

Individual control charts

Individual control charts also denoted to as Shewhart control charts are a form of SQC tool used to track and manage a process's variability over time. They are frequently employed in projects aimed at process and quality enhancement. A central line, which often represents the process means. The upper and lower control limits, which specify the process's allowable range of variation, make up an individual control chart. The process's data points are charted over time, and any points that deviate from the control range are considered to be out of control.

The methodological aspects are described below: For this study, the secondary data were collected from the public sector Central Pollution Control Board (CPCB), which is the official portal of the Government of India. The data contains more than 15 cities. It contains various air pollutant factors like PM2.5, PM10 and others, such as nitrogen dioxide (NO2), sulfur dioxide (SO2), carbon monoxide (CO), Ozone (ug/m3), RH (%), WS (m/s), WD (deg), SR (W/mt2), BP (mmHg), VWS (m/s) and other essential pollutants. Only information from the most recent three months (January to March 2023) was used for this investigation. Each component is measured every one hour.

RESULT AND DISCUSSION

This paper first examines the state of Tamil Nadu air pollution data with a particular emphasis on its cities. Based on the dataset's pollution levels, chose one district to focus on for our study in the beginning. The technique which is used takes into account several pollution indicators and measures to identify the most contaminated district. The average pollution of the Tamil Nadu state is shown in Fig. 2. The most polluted days in Tamil Nadu will be January 14–16, 2023, according to Fig. 2. Any events or functions might happen at that time. The objective of the paper is to select the areas with the worst pollution. Then we can determine which district has the worst pollution by examining the data on pollution for each district. Multiple pollutants including PM2.5 and PM 10, NO2, SO2, CO, and other pertinent pollutants are taken into account when calculating the most polluted district. Figure 3 shows the average polluted district in Tamil Nadu from the last three months data. After this procedure, we determined Tripur City as the most pollutant district. So we choose Tirupur city for further analysis. Then, as a statistical method, we employ PCA to find and prioritize the most important air pollution-related components. This algorithm has several steps. To make sure that all of the variables are on the same scale, we first standardize the pollutant data. The analysis cannot





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be dominated by variables with bigger degrees; hence standardization is essential. The covariance matrix is then computed using the standardized pollutant variables. The eigenvectors indicate the directions or patterns of variation in the pollutant data, and the eigenvalues represent the variance explained by each primary component. To find the primary components that best explain the variance, we order the eigen values in descending order. Each principal component's variance explanation shows how much of the total variability in the pollutant data that component was able to capture. We can also prioritize and concentrate our efforts on tackling the most significant pollutants or sources of pollution with the use of the PCA results (See Table 1 &Fig. 4), which are presented here. From Table 1 we conclude that PM 2.5 and PM 10 components of Tirupur city have significant impacts on air quality and are commonly monitored in air pollution studies. Descriptive statistics of PM2.5 and PM10 are given below in Table 2. There are significant data gaps in this report. Data that has been pre-processed by eliminating null values and substituting mean for the data. Python and Sigma XL software were used for this study.

Fitting of Individual control charts:

Control charts for people have been first recognised to determine the variation of the two variables PM 2.5 and PM10 components because the sample size utilised for the process monitoring is n=1. If the data are picked individually rather than in categories, the individual control charts are utilised to detect the presence of assignable causes as well as to establish the process level. Using the shifting range of two more observations, this type of graphic calculates the process variability. According to Fig. 5, both parameters are out of control since several points are outside the acceptable range and the oscillations of the points around the centreline are excessive and irregular.

Testing autocorrelation and trend

The control charts shown in Fig. 5 are based on the assumptions. First, we have to estimate the existence of autocorrelation. Fig. 6 represents the autocorrelations for the data. The straight lines are useful to estimate the non-zero correlation values of two standard deviation limits. (a) A bar extended outside the lines represents the statistical significance for both components. Since stationary series are required for the identification of the AR and MA components, it is important to check the data for trends before fitting the ARIMA model. ACF graphs in Fig. 6 are non-stationary because the autocorrelation is decreasing extremely slowly.

Fitting of ARIMA model

The final procedure model uses the difference in the d-th order transformation of the irregular variation in the "random walk" process to represent the stationary ARMA (p, q). In the ARMA (p, q) formula, the variables p and q represent the appropriate orders of the moving average and autoregressive components. To acquire an understanding, the PACF and ACF were examined. By utilizing substantial lags in ACF and PACF, we can select the significant ARIMA model. The procedures involved in creating an ARIMA control chart are the choice of a suitable model, estimation of the model parameters, and determination of control limits. The MSE and RMSE values of the model were used to choose the proper model. Less uncertainty in predicting mistakes is seen in improved ARIMA models' mean absolute error (MAE), mean squared error (RMSE), and mean absolute percentage error (MAPE). If the forecasts are accurate, the mean error (ME) and mean percentage error (MPE) should be very close to zero. The models chosen for the two components based on AIC values are listed in Table 3. For PM2.5 and PM10, respectively, ARIMA (1, 1, 3) and ARIMA (5, 1, 0) were defined as an appropriate models. The total amount of differences denoted as (d) for both models, since the data were differenced to provide stationarity. The resulting parameter estimates are shown in Table 4. A t-test and each estimated model coefficient are displayed. As a result of the coefficients' p-values being less than 0.05, the coefficients at the 5% level of significance are substantially different from 0.

Fitting of Special Cause Control charts(SCC)

Based on SCC tables for residuals The ARIMA model reveals that two PM2.5 and PM10 are beyond the set thresholds. Since some areas are outside of what can be controlled. This indicates that air pollution has not been under control for the last three months, and the cause of the change should be looked into. The IC control charts (Fig. 5) are unstable and uncontrollable. Traditional control charts demand that process observations be made in isolation from one another. The traditional control charts issue a high amount of false alarms when this criterion isn't met. The





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effectiveness of these graphics is significantly affected by even extremely minor levels of autocorrelation in the observations. SCC charts, the classic control charts used to observe the residuals of the ARIMA model; they are more efficient in this situations because they offer a higher probability of detection than individual charts.

CONCLUSION

In this paper, the highest pollutant cities were determined by using several pollutant measures and indicators. Tirupur is identified as the most pollutant city in Tamil Nadu. The next procedure using statistical tools like PCA to get the primary air components with the greatest amount of variance explained, sort the eigenvalues in descending order. The amount of variability that each principal component can capture is shown by the variance explained. As a result of PCA the air pollutants PM 2.5 and PM 10 are selected as the most significant pollutant for Tirupur District. Traditional control charts demanded that process observations be made in isolation from one another. The standard control charts issue a high number of false alerts when this requirement is not met. The performance of these charts is significantly impacted by even extremely low levels of autocorrelation in the observations. SCC charts, which are conventional control charts used to observe the ARIMA model's residuals, are more suitable related to environment since they give a better probability than individual charts. In this study, SCC charts for PM 2.5 and PM 10 are based on ARIMA (1, 1, 3) and ARIMA (5, 1, 0) models respectively. There are a few points out of control at the start of the second week of January and also at the March end. Compared to individual charts, residuals of fitted ARIMA model charts are valid for autocorrelation data. There must be many reasons like festivals, weekends, etc. To evaluate the performance and effectiveness of ARIMA control charts and individual control charts in monitoring and evaluating process stability, this study analysed and tracked the levels of air pollution in the Tirupur area. These findings demonstrated that the SCC chart, which offers a larger likelihood of coverage than individual control charts, is better appropriate for using auto-correlated data to assess the stability of the data. Even though it is essential to examine the assignable causes and try to recover the problem of the process. Quality of the air is important to the public health.

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Table 1: Eigen values and variance explained

Components	Eigenvalues	Variance
PM2.5 (ug/m3)	68416.1191	82.26%
PM10 (ug/m3)	9193.5183	11.05%
NO (ug/m3)	3698.6877	4.45%
NO2 (ug/m3)	1002.0801	1.20%
NOx (ppb)	669.8080	0.81%
SO2 (ug/m3)	126.0290	0.15%
CO (mg/m3)	46.0467	0.06%
Ozone (ug/m3)	19.0137	0.02%
RH (%)	0.4251	0.00%
WS (m/s)	0.2279	0.00%
WD (deg)	0.0000	0.00%
SR (W/mt2)	0.0000	0.00%





Table 2: Descriptive Statistics

	PM 2.5	PM 10	
Count	2160	2160	
Mean	38.3362	55.4382	
Std	18.5991	23.3344	
Min	4.3700	6.5800	
Max	196.0900	235.1700	
25%	25.9950	40.1220	
50%	37.0000	55.1850	
75%	47.6575	65.7900	

Table 3: Model comparison

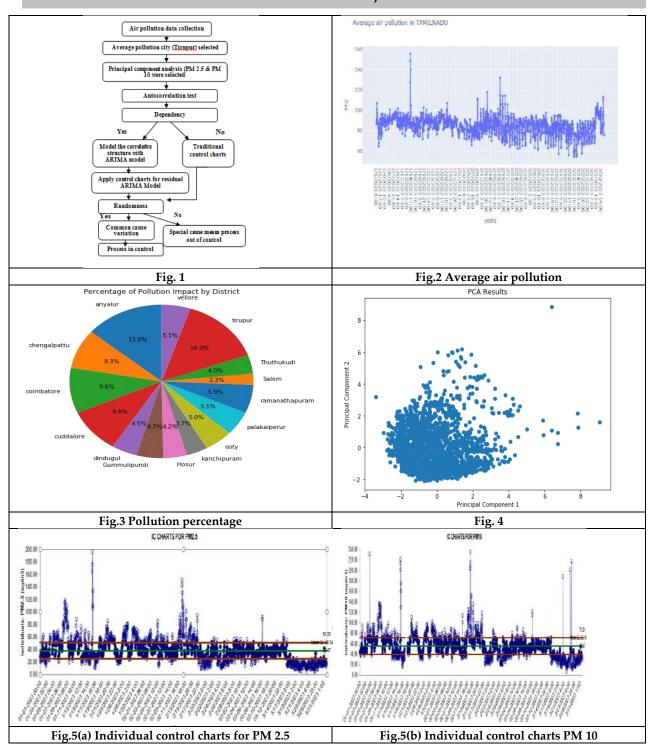
Table 3: Model comparison						
Parameter	Model	RMSE	MAE	MAPE	MASE	
	ARIMA(1,1,5)	8.0351	5.000	14.9541	1.0336	
	ARIMA(1,1,3)	8.0189	4.9573	14.7119	1.0247	
PM 2.5	ARIMA(4,1,1)	8.0283	4.9393	14.5958	1.0244	
	ARIMA(2,1,1)	8.1711	4.92317	14.5075	1.0174	
	ARIMA(3,1,1)	8.0407	4.9220	14.5856	1.01731	
	ARIMA(4,1,0)	14.0399	7.1577	13.5114	1.1995	
	ARIMA(5,1,0)	14.0279	7.1936	13.6025	1.2055	
PM 10	ARIMA(3,1,.0)	14.0492	7.0800	13.3737	1.1865	
	ARIMA(2,1,0)	14.0880	6.9796	13.1952	1.1697	
	ARIMA (1,1,0)	14.1338	6.7747	12.7704	1.9353	

Table 4: Parameter estimates

Parameter Model Model parameter		Estimate	Std Error	t	p	
		AR(1)	0.7030	0.09	7.71	0.00
		MA(1)	0.7510	0.103	7.265	0.00
PM2.5	ARIMA	MA(2)	0.3366	0.053	6.322	0.000
	(1,1,3)	MA(3)	-0.1483	0.077	1.910	0.05
		Constant	-0.0417	0.050	0.823	0.41
		AR (1)	-0.4183	0.043	9.572	0.00
		AR (2)	-0.3131	0.046	6.715	0.00
PM10	ARIMA	AR (3)	0.2457	0.047	5.181	0.00
PMH	(5,1,0)	AR (4)	-0.1846	0.046	3.964	0.00
		AR (5)	-0.0906	0.043	2.077	0.03
		Constant	-0.0863	0.1699	0.507	0.61

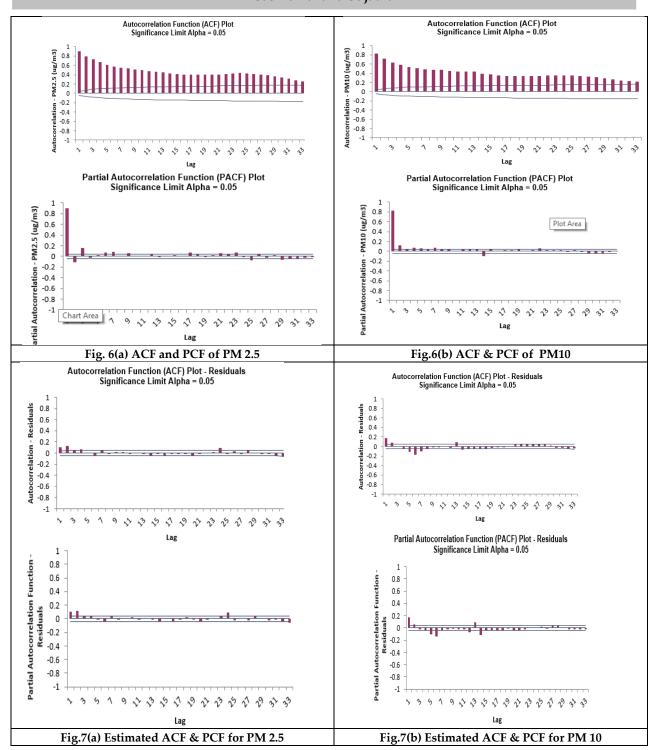






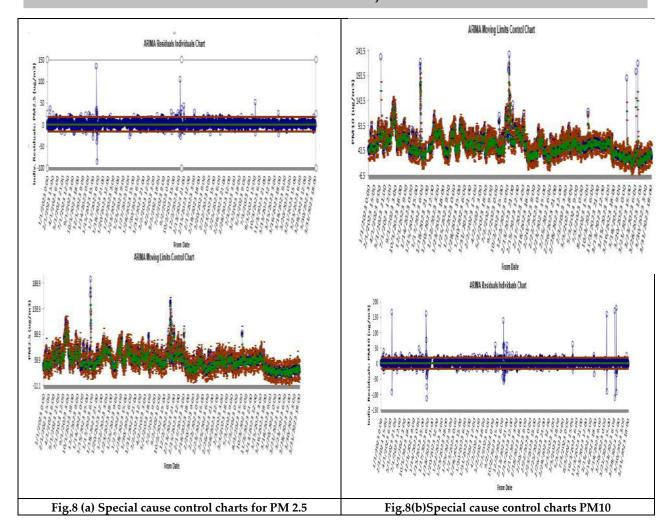
















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RESEARCH ARTICLE

The Significance of Visual Design Principles in Enhancing Data **Visualization: A Conceptual Perspective**

M. Jebakumar

Assistant Professor, Department of Visual Communication, St. Xavier's College (Autonomous), Palayamkottai – 627002, Tamil Nadu, India.

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*Address for Correspondence

M. Jebakumar

Assistant Professor, Department of Visual Communication, St. Xavier's College (Autonomous), Palayamkottai - 627002, Tamil Nadu, India. Email: jebakumarmp@gmail.com



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ABSTRACT

Data visualization plays a crucial role in effectively communicating complex information and facilitating decision-making. The effectiveness of data visualization largely depends on the application of visual design principles. This research paper explores the significance of visual design principles in enhancing data visualization from a conceptual perspective. Through a case study approach, the paper examines three reports from reputable organizations: UNICEF, WHO, and the World Bank. By employing Gestalt Theory of Visual Perception and Cognitive Load Theory as the theoretical framework, the study analyses how the application of design principles such as simplicity, consistency, colour and contrast, hierarchy, and organization contribute to the effectiveness of data visualization. The results demonstrate that adhering to these principles significantly enhances the clarity, readability, and comprehension of complex data. The findings highlight the importance of incorporating visual design principles in data visualization to effectively communicate critical information and drive positive change. This conceptual perspective underscores the need for data visualization practitioners to prioritize the application of visual design principles in their work to maximize the impact of their visualizations.

Keywords: Data Visualization, Visual Design Principles, Gestalt Theory, Cognitive Load Theory, Data Science, Visual Communication, Data Communication





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INTRODUCTION

Data visualization techniques are crucial in today's data-driven world due to the challenges posed by the vast amount of data being generated daily. These techniques enable individuals to quickly interpret and extract meaningful insights from complex datasets, facilitating faster and more accurate decision-making [1]. By choosing the right visualization techniques, such as charts, graphs, maps, and interactive dashboards, analysts can effectively convey desired insights. Visualizations also enhance comprehension and help in communicating insights to stakeholders, influencing them, and driving meaningful outcomes. Data visualization aids in data processing, identifying patterns, and developing ideas from large volumes of data. It enables better business understanding by presenting information in a visually appealing and informative manner[2]. Visual design principles play a crucial role in improving the effectiveness and understanding of data visualizations. These principles help in creating effective scientific visuals and improving the visual message. By utilizing principles such as hierarchy, contrast, alignment, and colour theory, data visualizations can be optimized for clarity, readability, and impact. These principles help in guiding the viewer's attention, simplifying complex information, and conveying insights more effectively[3]. In essence, the application of visual design principles is crucial for creating data visualizations that are not only visually appealing but also facilitate better comprehension and interpretation of the data presented. This research paper is to explore and understand the significance of visual design principles in enhancing data visualization from a conceptual perspective. By examining the various visual design principles and their application in data visualization, this study aims to shed light on how these principles contribute to the overall effectiveness and success of data visualization projects.

Review of the Literature

Visual design principles provide structured guidance on how to design compelling thumbnails for data stories and improve the overall effectiveness of data visualization. Kim et al. conducted a user study to understand the design choices for visualization thumbnails and found that different chart components play different roles in attracting reader attention and enhancing reader understandability[4]. They include determining the message before starting the visual, using appropriate colour combinations, and following established perceptual and cognitive principles[5]. Additionally, different chart components play different roles in attracting reader attention and enhancing reader understandability of visualization thumbnails. Design choices such as resizing, cropping, simplifying, and embellishing charts can make visualization thumbnails more inviting and interpretable[6]. Data visualization is the graphical representation of information and data, using visual elements like charts, graphs, and maps to communicate complex data clearly and efficiently. It allows for quick decision-making and is crucial for businesses, researchers, and analysts when conveying their findings. By presenting data visually, it becomes easier for the audience to understand and interpret the underlying trends, patterns, and insights. Visualizations turn raw data into a story, making it easier to identify and understand significant information. Murchie and Diomede discuss the need or guidance on improving the visual aspects of science communication, including data visualization [7]. Kelleher and Wagener emphasize the need for clear and concise visual representations of data, stating that well-designed visualizations can facilitate understanding and decision-making. They propose a set of guidelines for creating effective data visualizations, including the use of appropriate visual encodings, the consideration of the target audience, and the reduction of clutter [8]. Sainz Sujet emphasizes the importance of understanding the principles of data visualization for elegant and efficient design [9]. Few argues that the application of visual design principles is crucial for creating compelling and informative data visualizations. He suggests that effective visualizations should be aesthetically pleasing, easy to understand, and focused on conveying the most important information [10]. According to Fernandes and Steele, visual design principles play a vital role in creating effective data visualizations. They proposed that principles such as simplicity, coherence, and colour contrast contribute significantly to the clarity and understanding of the data being visualized. In their study, the authors demonstrated that adhering to these principles can improve the accuracy and speed of decision-making based on the visualized data [11]. Another critical aspect of visual design principles in data visualization is the use of Gestalt principles. As suggested by Nesbitt and Friedrich, Gestalt principles such as proximity, similarity, and continuation can be employed to guide the viewer's





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attention and create a more coherent and organized visualization [12]. Harrower and Brewer proposed a colour scheme selection tool for data visualization, emphasizing the importance of colour consistency and contrast for effective visualization. They argued that appropriate colour schemes can significantly enhance the readability and understandability of data visualizations[13]. Balance and contrast are foundational design principles that contribute to the stability and clarity of visual presentations. Borkin et al. found that balanced compositions tend to be perceived as more professional and trustworthy, which is crucial for the credibility of data visualizations [14].

MATERIALS AND METHODS

The purpose of this paper is to conceptually examine and highlight the importance of applying established visual design principles to enhance the effectiveness of data visualizations. Specifically, it aims to review key visual design theories and principles, analyse how they facilitate extraction and comprehension of insights from data visualizations, and discuss their significance in the context of the growing reliance on data visualization across domains.

This research employed a case study approach to investigate the significance of visual design principles in enhancing data visualization from a conceptual perspective. Case studies were chosen as they allow for an in-depth examination of specific instances, providing rich insights into the application of visual design principles in data visualization[15]. Three case studies were selected from World Wide Data Visualization Reports published by reputable organizations: the World Health Organization (WHO), the World Bank, and the United Nations International Children's Emergency Fund (UNICEF). These reports were chosen due to their extensive use of data visualization techniques to communicate complex information effectively.

This research employed Gestalt Theory of Visual Perception and Cognitive Load Theory. Gestalt Theory explores how people organize visual information into meaningful patterns and relationships[16]. Key principles such as figure-ground, similarity, proximity, closure, and continuity provide insight into how viewers interpret visual designs. These principles should inform choices made in data visualization regarding layout, colour, shapes, etc [17]. The Cognitive Load theory suggests that there are limits to how much information can be processed effectively by the human brain at one time[18]. When these limits are exceeded, it can lead to cognitive overload, which hinders learning and retention. The integration of both theories provides a robust theoretical framework for examining the significance of visual design principles in enhancing data visualization.

RESULTS AND DISCUSSION

Case study 1: UNICEF Children Migration Report 2016

This report incorporates a diverse range of data visualization techniques to present statistics, trends, and narratives related to refugee and migrant children[19]. The report includes charts, graphs, maps, and infographics to present data on population demographics, education, health, and protection issues (see Fig. 1). These visual aids effectively support the narrative and provide a comprehensive understanding of the challenges faced by these vulnerable children. The application of design principles significantly enhances the effectiveness of data visualization in the UNICEF report. The following design principles were observed: The report utilizes a simple and clean design, ensuring that the visualizations are easily comprehensible. Complex data is simplified through the use of clear and concise visuals, making it accessible to a wide audience. The use of colour and contrast in the visualizations helps to highlight key information and draw attention to important data points. The colour palette is carefully chosen to create visual impact and facilitate quick understanding. The report effectively employs hierarchy and organization in its visual design to prioritize and structure information. This allows the audience to navigate through the content seamlessly and comprehend the data in a logical manner (See Fig. 2). The effective use of data visualization techniques, along with design principles such as simplicity, consistency, colour and contrast, hierarchy and organization aids in communicating complex information in a clear and engaging manner. This conceptual





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perspective highlights the importance of incorporating visual design principles in data visualization to effectively communicate critical issues and drive positive change.

Case Study 2: WHO - World health statistics 2023

Data visualization in the WHO's report is primarily presented through graphs, charts, and infographics[20]. These visualizations effectively communicate global health trends and the progress of the SDGs. For instance, the use of line graphs to depict changes in health indicators over time, bar charts to compare health statistics across regions, and world maps to show geographical disparities in health outcomes (see Fig. 3). The design principles employed in the WHO's report include simplicity, clarity, consistency, and relevance. The report maintains a simple layout, which enhances readability and understanding. The use of clear, legible fonts and appropriate colour schemes contributes to the clarity of the report. Consistency in the use of visual elements, such as the same scales for axes in graphs, helps in easy comparison across different data sets. The relevance of the visuals to the data and the message they convey is also noteworthy (see Fig. 4). The results show that the effective use of design principles significantly enhances the data visualization in the WHO's report. Clear and simple visuals make it easier for readers to understand complex health data. Consistency in visual elements allows for easy comparison between different data sets. The relevance of the visuals to the data and message ensures that the intended information is effectively communicated.

Case study 3: World Bank – World Development Report 2023

This report effectively employs data visualization to convey complex information about migration and its impact on societies[21]. The report uses a variety of charts, graphs, and maps to illustrate key findings, such as:Global migration patterns and trends. The distribution of refugees and displaced persons, the economic and social impacts of migration on both sending and receiving countries (see Fig 5). Visualizations that adhere to these principles are more likely to communicate insights accurately and persuasively. Bar Chat (see Fig. 6) illustrating migration flows are more comprehensible when using consistent colour coding and appropriate legends. Similarly, charts and graphs that prioritize clarity and simplicity facilitate quicker comprehension of complex data relationships. The results demonstrate that well-designed data visualizations can effectively distil complex information related to migration and refugee movements into accessible and actionable insights. Through strategic application of design principles and the visualizations featured in the report succeed in conveying key messages and facilitating informed decision-making.

CONCLUSION

This research paper has explored the significance of visual design principles in enhancing data visualization from a conceptual perspective. Through a comprehensive review of the literature and an in-depth analysis of three case studies, it has become evident that the application of visual design principles plays a crucial role in creating effective and impactful data visualizations. The case studies examined in this paper, namely the UNICEF Children Migration Report 2016, the WHO World Health Statistics 2023, and the World Bank World Development Report 2023, have demonstrated how the strategic use of visual design principles can significantly improve the clarity, readability, and persuasiveness of data visualizations. By employing principles such as simplicity, consistency, colour and contrast, hierarchy, and organization, these reports have successfully communicated complex information in an accessible and engaging manner.

The findings of this study have several implications for the field of data visualization. Firstly, they underscore the importance of incorporating visual design principles into the creation of data visualizations, as they can greatly enhance the effectiveness and impact of the visualizations. Secondly, the results suggest that a thorough understanding of visual design principles is essential for professionals working in the field of data visualization, as it enables them to create visualizations that are not only aesthetically pleasing but also highly functional and informative. The integration of Gestalt Theory of Visual Perception and Cognitive Load Theory provides a robust theoretical framework for understanding the significance of visual design principles in data visualization. Gestalt





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Theory highlights the importance of organizing visual information into meaningful patterns and relationships, while Cognitive Load Theory emphasizes the need to present information in a way that does not overwhelm the viewer's cognitive processing capacity. Moreover, this research highlights the need for further exploration and development of visual design principles specifically tailored to the field of data visualization. As the amount and complexity of data continue to grow, it is crucial to establish a set of best practices and guidelines that can help practitioners create visualizations that are both visually appealing and intellectually stimulating.

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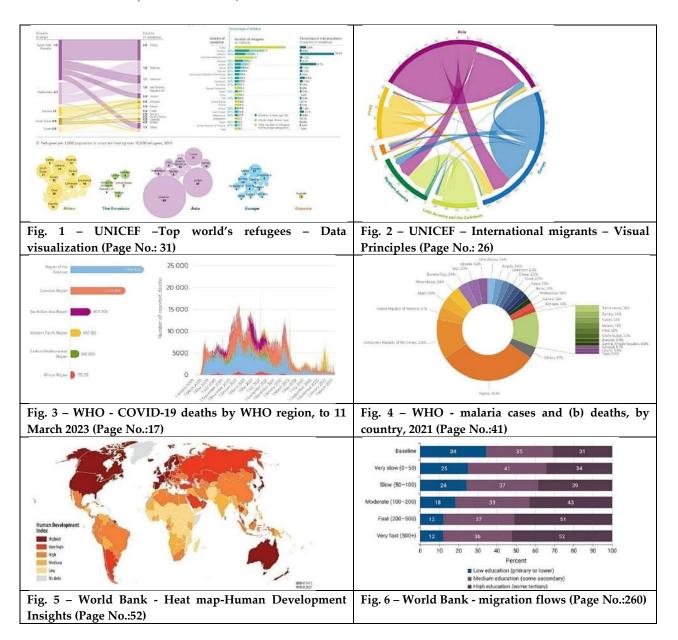
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RESEARCH ARTICLE

Investigations of Efficient Nonlinear Optical Tetra Glycine Ammonium Sulphate (TGAS) Crystal for Optoelectronic Applications

Josephine Gladiya M^{1a}, Anuradha G V², Sivashankar V^{1*}, Helina B¹

^aResearch Scholar, 20211282132008

Department of Physics, St. Xavier's College (Autonomous), Palayamkottai - 627 002, Tirunelveli, Tamilnadu, India.

²Department of Physics, Sri Parasakthi college for Women (Autonomous), Courtrallam, Tenkasi – 627 802, Tamilnadu, India

Affiliated to Manonmaniam Sundaranar University, Tirunelveli - 627 012, Tamilnadu, India

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*Address for Correspondence Sivashankar V

Department of Physics, St. Xavier's College (Autonomous), Palayamkottai – 627 002, Tirunelveli, Tamilnadu, India. Email: tvsivashankar@yahoo.co.in



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ABSTRACT

Current study focuses on synthesized Tetra glycine ammonium sulphate crystals (TGAS), which is grown by traditional solution growth approach and its unique properties. Single crystal as well as Powder XRD confirm the orthorhombic pattern of TGAS crystal. The lower cut off is 199 nm for TGAS crystal has been obtained from UV – Vis spectra and the crystal exhibit greater transparency in the visible zone. The high band gap value of 6.25 eV has been enumerated using Tauc's plot. The luminescent property has been confirmed by PL spectrum. The hardness and work hardening coefficient of 2.642 of TGAS crystal was obtained from Vickers hardness analysis. The TGAS crystal withstands the higher temperature of 240 °C.Using EDX analysis, every element found in the TGAS crystal in the predicted amounts was detected. The NLO property was confirmed by SHG analysis and the crystalline sample of TGAS is 1.17 time greater than KDP and the high Laser damage threshold of 2. 04 GW/ cm² has been obtained for the TGAS crystal. The enhanced and unique features of structural, optical, thermal, mechanical properties with greater SHG efficiency and Laser damage threshold for the TGAS crystal emphasize that TGAS is a superior candidate for fabricating Optoelectronic gadgets

Keywords: TGAS crystal, NLO, SHG, XRD





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INTRODUCTION

Accentuately, Amino acids are the greatest torchbearers of crystal growth, as they're the organic compounds that comprises the astonishing optical, thermal and mechanical properties with greater Second Harmonic Generation (SHG) efficiency [1]. The quest for novel materials with enhanced characteristics is the leading lights of today's science and technology [2]. The new organic materials with wide transparency domain, greater thermal stability, higher mechanical hardness, relatively good Second harmonic generation efficiency is the desperate necessity in academic research as well as in technological aspects [3]. In order to grow such novel materials with meticulous properties, Researchers used commercially available α – Glycine which is a stable and simplest of all amino acids [4,5]. Glycine is the Researcher's ecstasy, as it amuses the investigators with its eccentric and noteworthy characteristics [6]. Glycine is a non-polar achiral molecule, that can extensively incorporate with all the organic and inorganic compounds to produce novel materials with astonishing physiochemical properties [7]. Researchers have grown various glycine crystals utilizing organic and inorganic compounds, some are listed as follows: Gamma glycine crystals from cerous chloride [7], lithium nitrate [8], Manganese sulphate [9],Di glycine crystals [10], Tri glycine crystals [11] and Glycine doped crystals [12] have been reported and their unique properties have been scrutinized. Principal objective of this manuscript is to investigate and provide illumination to the novel material of Tetra Glycine Ammonium sulphate (TGAS) single crystals and their astonishing properties are discussed for the first time. The single crystalline Tetra Glycine Ammonium Sulphate are(TGAS) grown by adapting conventional solution growth approach at ambient temperature. The various structural, spectral, mechanical, thermal, electrical and elemental characteristics of the proposed TGAS crystal have been briefly discussed.

Experimental Procedure

Single crystals of Tetra Glycine ammonium sulphate are grown by dissolving Glycine (Merck) of purity >=99% and Ammonium sulphate (Hi − media) of purity ≥99%in aa 4:1 ratio added to 100 milliliters of deionized water in a beaker. After that, the solutions in the vessel were agitated at a consistent temperature of 50 degrees for the next five hours. Following a Whatman filter paper filtering the saturated solution, the beaker was capped with perforated aluminum foil and allowed to slowly evaporate at room temperature. In less than 30 days, high-quality single crystals of Tetra Glycine Ammonium Sulphate (TGAS) were obtained. Theharvested crystal of TGAS is depicted in Fig: 1

Characterization

To ascertain the lattice specifications and the structure of the TGAS crystal, a Bruker D8 Quest X-ray diffractometer of MoK $_{\alpha}$ radiation (λ = 0.710 Å) was employed. The crystal of TGAS was scrutinized bypowder XRD diffractometer Bruker AXS D8 Advance (PANalytical, Nickel filtered Cu K $_{\alpha}$ (λ = 1.540 Å) at 35 KV, 10mA). The optical transmission spectra for the TGAS crystal were collected viavarian Cary 5000 spectrometer in the region 190-1100 nm at room temperature. PL spectrum and it was recorded by photoluminescence spectrometer, Fluorolog HORIBA in 200 to 900 nm range with the wavelength of excitation about 250 nm. For hardness analysis, a diamond indenter-equipped Shimadzu Vickers microhardness tester connected to an incident light microscope was utilized. The thermal endurance of TGAS crystal was found by carrying out TG/DTA plot using a NETZSCH STA 409C/CD thermal analyzer with nitrogen atmosphere in 30-400°Ctemperature at the heating rate 10°C/min. The spectrum of EDX is collected employing a high-resolution scanning electron microscope (HRSEM), Thermo Scientific Apero S. A Q-switched Nd: YAG laser of model HG-4B was used with the repetition rate of 10 Hz and pulse width 6 ns in the Kurtz-Perry powder method. Using a high intensity Q-switched Nd: YAG laser with wavelength of 1064 nm and a pulse width of 10 ns, the LDT value of the grown TGAS crystal was determined. The antibacterial activity study is used to check the grown TGAS crystal's resistance to bacterial specimens.





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RESULTS AND DISCUSSIONS

Single crystal XRD

Table 1 displays the acquired lattice specifications for the TGAS crystal. From the specifics, it is evident that the TGAS crystalline material crystallizes under orthorhombic system. The lattice specifications of previously reported Glycine ammonium sulphate crystals are compared with the parameters obtained for TGAS crystal are presented in table 1. It is apparent that the Tetra glycine ammonium sulphate crystal varies from previously reported Glycine ammonium sulphate crystal which crystallizes in a monoclinic system whereas TGAS crystallizes in an orthorhombic system.

PXRD Studies

The powder XRD spectrum of TGAS is illustrated in Fig.2. Utilizing the WinXMorph software, the corresponding Miller indices (hkl) were assigned for the TGAS crystal. The intense sharp diffraction peaks at $20 = 19.025^{\circ}$ and 25.293°indicates the (110) and (013) planes and medium intense peaks appears at $20 = 14.815^{\circ}$, 23.002°, 23.954°, 30.232°, 31.126°, 33.624°, 36.594°, 42.548°, 53.422°, 63.299° belongs to (002), (111), (112), (113), (022), (121), (212), (304) and (135) planes respectively that confirms the existence of orthorhombic crystalline structure. The remarkable diffraction peaks reveal the exceptional crystallinity, high purity and good structural precision of the TGAS crystal [14]

FTIR Spectrum

Fig: 3 illustrates the FTIR spectra for the TGAS crystal. The intense and wide absorption peak at 3418 cm⁻¹denotes the OH stretching [15]. The peak at 2612 cm⁻¹ belongs to C=O stretch [16]. The peak at 2229 cm⁻¹ denotes the intense sulphate overtones and combinations [17]. The peak at 2173 cm⁻¹ and 2057 cm⁻¹belongs to the CO stretching [18,19]. The peak at 1630 cm⁻¹denotes OH stretching [20] and the peak at 1590 cm⁻¹ confirms COO ie, the stretching of carboxyl group [21]. The peak at 1488 cm⁻¹ indicates the C=O vibration [22]. The peak at 1404 cm⁻¹ indicates C-N vibration [23]. The peak at 1334 cm⁻¹ indicates the in-plane bending of CH group [24]. The peaks at 1272 cm⁻¹ and 1127 cm⁻¹ indicates C-O band [25] and C-O-C stretching vibration [26]. The peak at 1041 cm⁻¹ and 929 cm⁻¹ indicates the presence of C-O stretching [27] and wagging vibration of OH group [28]. The peaks at 889 cm⁻¹ and 833 cm⁻¹ indicates the presence of C-O stretching [29] and OH stretching [30]. The peak at 685 cm⁻¹ belongs to the torsion in hydrogen bonds of amide [31]. The peaks at 609 cm⁻¹ and 503 cm⁻¹ indicates the presence of OH bending vibration [32] and C-H out of plane bending vibration [33].

UV – Vis spectrum

The transmittance as well as absorbance spectra of TGAS crystal are displayed in the fig:4 and 5. From the plot, it is evident that the crystal possesses less absorbance in the complete UV, visible and near infrared zone. The lower cutoff for TGAS crystal is 199 nm. It is apparent that the transmittance spectrum of TGAS crystal shows greater transparency in the entire visible zone. The optical band gap was scrutinized from the Tauc's plot as illustrated in Fig.6. The band gap energy E_8 of TGAS crystal is obtained to be 6.25 eV which is greater than the previously reported undoped Ammonium Sulphate which is 5.6 eV[34]

Photoluminescence Analysis

Fig 7 illustrates the PL spectrum for the TGAS crystal. The peaks of emission at 462 nm, 540 nm are due to the emission of visible light. The emission peak at 822 nm belongs to the IR light emission. The intense peak patterns reveal that the TGAS crystal has high ordered crystallinity, utmost purity and structural reliability, implying that TGAS crystal is a desirable luminescent material. The attainable PL emission conveys the crystal is an ideal choice for NLO applications[35].





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Hardness Studies

The micro hardness number was enumerated and the hardness number variation (H_v) with the (P) applied load of TGAS crystal is illustrated in the fig:8 and hardness gradually rises with the rise in applied load. The work hardening efficacy (n) was enumerated using the Meyer's relation and the value of n is found to be 2.462 from the Fig.9. Since n is greater than 1.6, the proposed TGAS crystal is a softer medium [36].

Thermal Analysis

Fig: 10depicts the TG plot for the TGAS crystal. The TG curve depicts there is no change in weight upto 230 °C. The first major weight loss of about 27.02% from 230 °C to 275 °C belongs to the sublimation and initial decomposition of TGAS crystal. The second weight loss of about 21.25% from 275 °C to 290 °C is due to further decomposition of TGAS crystal. The remaining sample melts and decomposes giving a weight loss percentage of about 17.05% from 290 °C to 400 °C. The DTA curve for the TGAS crystal is depicted in fig11. The DTA exhibits an intense endothermic peak at 240 °C which is the decomposition point for the TGAS crystal. The outcomes suggests that the TGAS has a greater thermal stability of about 240 °C which is higher than the previously reported Gamma glycine crystal of ammonium sulphate as solvent [37], and it is apparent that the TGAS crystal can be effectively applied for fabrication of Optoelectronic devices[38].

EDX Spectrum

Fig11 illustrates the recorded Energy Dispersive X ray spectrum for the TGAS crystal. In the plot, S C, O and N peaks subsequently located at 0.1, 0.15, 0,2 and 0.25 keV. For the TGAS crystal, the identified elements viz, C, N, O and S are in expected proportions as their atomic distributions of C, N, O and S are 35.23%, 13.48%, 51.13% and 0.16% respectively. Hence, it is obvious that there is no detection of foreign element in the EDX spectrum of TGAS crystal, that implies that TGAS is a high purity crystal and can be utilized for fabricating Nonlinear optical devices[39].

Second Harmonic Generation Analysis

For measuring SHG efficacy, KDP was taken as a reference material. The incident wavelength is 1064 nm with emitted wavelength of light from the crystal is 532 nm (green laser light). The input energy is 0.5 J for both KDP and TGAS sample. The output energy is 8.75mJ for TGAS crystal and it is 7.5 mJ for KDP. Hence, the crystal of TGAS is 1.17 times that of KDP sample and it is greater than the previously reported Glycine ammonium sulphate crystals (0.90 times that of KDP)[13], since the SHG efficacy of TGAS crystal is high, it can be utilized for NLO applications[40]. Table 2 shows the obtained value of SHG efficacy for TGAS crystal.

Laser Damage Threshold studies

The equation for estimating the LDT of a TGAS crystal is Power density (P)= $E/t\pi r^2$, and the LDT value for the grown TGAS crystal was enumerated to be 2.04 GW/cm². Since it is more than LDT of KDP crystal (0.20) and Urea (1.50) [41] TGAS could be used for laser applications.

Antibacterial Activity

Antibacterial activity was carried out and the inhibition data were evaluated for the produced crystal of TGAS with the goal to obtain information about the bacteria and other microorganisms that tend to proliferate on the outermost layer of the crystal [42]. The examination of the antibacterial properties of the TGAS crystal involves two bacterial specimens viz., *Staphylococcus aureus* and *Pseudomonas aeruginosa*. The zones of inhibition areas for the grown crystal of TGAS and the standard sample are given in the figure 10. The data pertaining to the investigation of antibacterial activity is specified in the table 3. Considering the outcomes, it is apparent that the accomplished crystal of TGAS has lesser inhibitory effect on bacterial specimens, and it is therefore inferred that the crystal has less effective antibacterial properties [43]





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CONCLUSION

Tetra glycine ammonium sulphate crystal was synthesized by utilizing Solution growth approach. The harvested crystal was examined with various spectral, structural, optical, thermal, mechanical, elemental, SHG and LDT analyses. Both the Single crystal and Powder XRD validates the presence of orthorhombic crystalline structure. UV-Vis spectra confirms the TGAS crystal has greater transparency in the complete visible zone with higher band gap value of 6.25 eV. Photoluminescence studies reveals that the TGAS crystal exhibits good luminescence. Hardness analysis reveals that the TGAS crystal is a softer medium. Thermal analysis conveys that the TGAS crystal possesses higher thermal durability of 240 °C. EDX spectra reveals that the TGAS crystal is a high purity material. The TGAS crystal possesses greater Second Harmonic generation efficacy of 1.17 times greater than KDP with greater LDT of 2.04 GW/ cm². The TGAS crystal is found to have less antibacterial activity. Thus, the TGAS crystal is found to exhibit greater transparency, higher band gap value, good structural perfection, high purity, good luminescence, greater mechanical and thermal stability, higher NLO efficiency and greater Laser damage threshold values, it could potentially be feasible to design optoelectronic gadgets utilizing the TGAS crystal.

Research Data Policy and Data Availability

The datasets generated or analyzed during the current study are included in this submitted article.

Author Contribution

Josephine Gladiya M: Conceptualization, methodology and writing - Original draft,

Anuradha G. V: Validation, Investigation and review,

Sivashankar V: Correction proof, Validation and review.

Helina B: review

All authors read and approved the final manuscript.

Ethics Declaration

Consent to Participate

All authors have given consent to participate

Funding and Competing Interest

The authors did not receive support from any organization for the submitted work. The authors declare no competing interests.

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Table 1: SXRD data for TGAS crystal

Parameters	SXRD data for TGAS crystal	SXRD data for GAS
		Crystal
Crystal structure	Orthorhombic	Monoclinic
A	90°	90°
В	90°	92. 66°
Γ	90°	90°
a(Å)	11.981(1) Å	8.262(3) Å
b(Å)	21.234(3) Å	10.074(2) Å
c(Å)	7.784(2) Å	8.632(2) Å
V(Å ³)	1980.28(2)Å ³	717.39(4) Å ³
References	Present work	[13]

Table 2: Second harmonic efficiency data for TGAS crystal

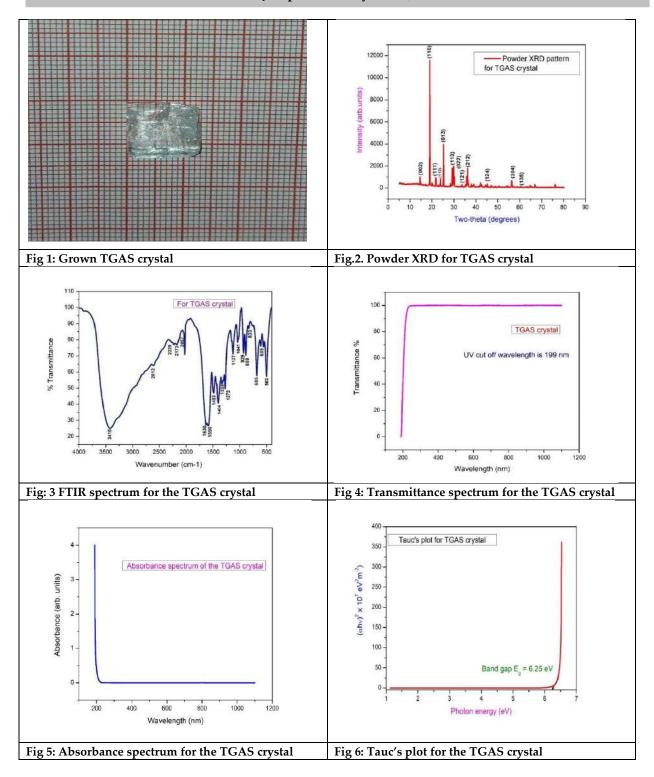
S. No		Output Energy	Input Energy
	Name of the Sample	(milli joule)	(joule)
1	KDP (Reference)	7.5	0.50
2	TGAS crystal	8.75	0.50

Table 3:Data for Antibacterial activity of TGAS crystalline sample

Name of the Bacteria	Zones of inhibition in TGAS crystal (mm)	Standard values of Zones of inhibitions (mm)
Staphylococcus aureus	20	25
Pseudomonas aeruginosa	15	18

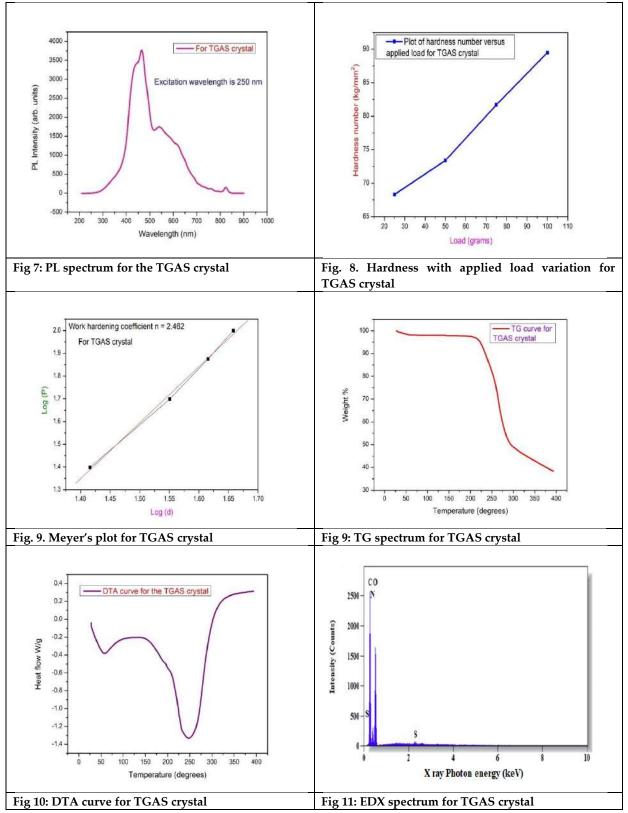
















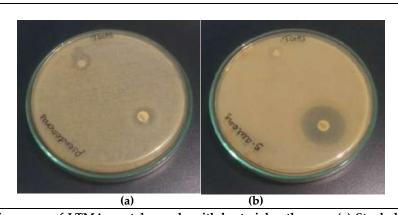


Fig 12:The inhibition areas of LTMA crystal sample with bacterial pathogens: (a) Staphylococcus aureus (b) Pseudomonas aeruginosa





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RESEARCH ARTICLE

Impact of Artificial Intelligence in Automobile Industry in India

K.Kevin Selva^{1*} and Dr.A.Natarajan²

¹Student, PG Department of Data Science, St. Xavier's college, Palayamkottai – 627 002, TamilNadu, India. ²Assistant Professor, PG Department of Data Science, St. Xavier's college, Palayamkottai – 627 002, Tamil Nadu, India.

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*Address for Correspondence K.Kevin Selva,

Ist M.sc Data science, Department of Data Science, St. Xavier's college, Palayamkottai – 627 002, Tamil Nadu, India. Email: kevinselva2000@gmail.com



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ABSTRACT

Artificial Intelligence (AI) is transforming a number of industries globally, and it has a significant and wide-ranging effect on the Indian automotive industry. Focusing on how AI is changing the Indian automotive industry, emphasizing the major areas in which AI technology is changing the scene. In this article we mostly discussed the consumer's interest in need of Autonomous Emergency Braking (AEB) and Driver Assistance System (ADAS)and other artificial intelligence system in Low/Mid-Range Cars and Other Automobiles for Safety and Business Reasons. Furthermore, AI is essential to India's development of connected and electric automobiles. Collecting data to give producers useful insights by analyzing industry trends and forecasting consumer preferences .In conclusion, the impact of AI on the automobile industry in India is profound and far-reaching, revolutionizing every aspect of the automotive value chain from design and manufacturing to sales and customer experience. As AI continues to evolve, it will be imperative for Indian automotive companies to embrace and harness its transformative potential to stay competitive in the rapidly evolving global automotive market.

Keywords: Artificial Intelligence, Automobile Industry, ADAS, AEB, Scope, Market trend, analysis.

INTRODUCTION

Artificial Intelligence(AI) is an alternative for human Intelligence in machines that are lined up to think like humans and implement the exact grasping techniques of human brain over various domain. AI encompasses the development of computer systems capable of performing tasks that typically require human intelligence, including learning, reasoning, problem-solving, perception, and decision-making[1]. AI serves for many purposes in a wide range of





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sectors and fields since it can carry out operations that would typically require human intelligence Automotive industry is one of the biggest investors in AI. It is propelling a significant shift in how the auto industry will develop in the future. Players in the automotive sector are enhancing the quality of data essential for autonomous driving systems by utilizing machine learning techniques. This guarantees more accuracy and safety in the operation of selfdriving cars. Technologies like ADAS monitors the area surrounding a car using a variety of technologies, including radar, cameras, and sensors. In considering this, ADAS gives drivers alerts and information in real time to help them steer clear of potential hazards like crashes. Depending on the ADAS tech's proficiency, the system can either fully take over control of the car to assist avoid any accidents, or it can alert the driver by sound, vibration, and display indications.[4] ADAS technology has up to five levels. Autonomous Emergency Braking (AEB) is defined as a system that constantly keeps track of the road ahead and will automatically halt the vehicle if the driver fails to take action. AEB works autonomously, with no input from the driver. The system can also add braking force if you are pressing the brake, but weaker than a vehicle needs to avoid a collision. Every AEB system monitors vehicles and most of the time, pedestrians and other obstacles. The goal of the AEB system is to avert or lessen serious crashes by applying the brakes when sudden dangers arise or if the driver is not taking appropriate action quickly enough and many other AI features can help towards safer and comfort driving experience and could make buyers towards AI integrated vehicles[5]. The growth of the Indian car market is encouraging. Owning a car is far more appealing to people than taking public transportation. Vehicle becomes sentiment for Indians as it results in Heavy traffic and pave way for accidents and loss of life. AI features can either recognize a threat to the car and engage the driver in order to prevent it from happening, or they can work automatically to steer the vehicle out of danger. When compared to low and mid-range vehicles, luxury cars are scarce in the Indian market.AI elements must be added to low- and mid-range vehicles in order to enhance passenger safety and satisfaction. Therefore, it is critical for manufacturers to understand that the Indian market will favor any vehicle with AI features that is within the affordability of the middle class. This article proves that buyers are interested and also ready to invest in artificial intelligence system in their cars and gives a prediction about the taste of future buyers.

Method Used For Analysis

The integration of Artificial Intelligence (AI) in the automotive sector has witnessed remarkable advancements, revolutionizing the industry's landscape. This article aims to provide a comprehensive overview of AI applications in automobiles, with a specific focus on leveraging gathered data for market analysis. In line with AI integration, the automobile sector is placing more and more focus on using collected data for insightful market analysis. A paradigm change has occurred with the advent of big data, allowing practitioners and scholars to identify important patterns and trends from enormous datasets. The convergence of AI and extensive datasets is driving transformative insights, shaping the future of the automotive industry. To obtain insightful information, a survey was conducted using Google Forms, and the raw data was gathered, cleaned, processed, visualized, and analyzed in Microsoft Excel.

Forecasting The Trend Of The Indian Automobile Market In Cars

The automobile market in India is one of the largest and fastest-growing in the world, playing a significant role in the country's economic development. With a diverse consumer base and a steadily increasing middle-class population, India has emerged as a crucial market for both domestic and international automotive manufacturers.

Market Overview

The Indian automobile market is characterized by a wide range of vehicles, including two-wheelers, four-wheelers, commercial vehicles, and electric vehicles, each contributing to the overall dynamism of the industry. The automobile Industry in India is a dynamic and evolving industry driven by consumer preferences, technological advancements, and government policies. As the country continues to urbanize and experience economic growth, the automotive sector is expected to play a crucial role in shaping India's mobility landscape.





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Visualizing the past 20-Years Automobile(CARS) sales

Over the last 20 years, there has been an increase in car sales. The graph shows statistics from 2004 to 2023. The car business is developing far faster than expected, as seen by the jump from one lakh cars sold annually to about 40 lakh sales. Fig 1

India's Global Automobile Economy Share

India is a prominent player in the global automobile market, but compared to certain other big economies, its economic share in terms of car sales is not as high. A growing middle class, more urbanization, and expanding purchasing power are some of the reasons driving the steady growth of the Indian automotive market It's important to remember, nevertheless that India's economic contribution to global auto sales is not as significant as that of countries like China, USA, or the EU. The largest automobile market in the world, in terms of both sales and production, has been China at the end of last decade. Fig 2

Report on India's Top 10 Automobiles

As of January 2022, the list of the top-selling cars in the Indian market can change over time based on consumer preferences, new launches, and market trends. Here's a general overview of some popular and consistently well-selling cars in India. The specific rankings and models might have changed since then. Fig 3

BEV and Hybrid vehicle's performance in Indian market:

Sales of BEVs and hybrid automobiles have seen ups and downs, with a primary cause being consumers' confusion when selecting a vehicle type. The chart clearly showcases the sales fluctuations on BEV and Hybrid sales of 2023 market Fig 4

Data Analysis - Understanding Buyer's Mindset

This article intend to close the gap between buyers and manufacturers in this particular component. As seen previously, the market as a whole is expanding, but consumers are still having trouble selecting a car. Data from present and potential consumers to learn about their preferences and mindsets. Also to know if they were genuinely interested in having artificial intelligence (AI) systems installed in their vehicles.

List of questions placed to the audience and reason for the questions:

	of questions placed to the audience and	1
No.	Questions	Explanation
1.	Type of participant	Understanding their role is crucial in order to comprehend their
	Student	attitude
	Working professional	
	Business and others	
2.	Does the individual currently possess	Future vehicle buyers may have different ideas about what kind of
	a car?	car to buy than someone who already owns one.
	• Yes	
	• No	
3.	The type of car model they would like	The world is shifting towards electric vehicles (EVs), therefore we
	to purchase in the future	can analyze whether people are prepared to prefer EVs and hybrids
	Petrol/Diesel	over traditional gasoline and diesel vehicles.
	Hybrid	
	• BEV	
4.	Preference of price range when they	The primary goal of this study is to apply artificial intelligence to
	buy in future	low- and mid-range vehicles. By asking this question, we can
	• Low(Upto 6L)	ascertain what price range consumers are willing to invest in.
	• Mid(6-15L)	
	High(Above 15L)	
5.	Interested in installing AI feature in	After outlining the features of AI, participants are asked if they





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	your car(Features like ADAS, ABS etc)	would want to have it in their cars. This helps us to understand the
	• Yes	participant's interest in AI
	• No	
6.	What will you prefer?	To determine whether individuals are interested in investing for
	A car with AI features with a	Artificial Intelligence
	lakh higher than its actual price	-
	A car without AI feature	

Examining the survey's findings:

The purpose of the survey is to determine consumer interest in AI integration into automobiles as well as the considerations that automakers should make when developing new models or making adjustments to current models. Fig 6

Types of People Participated

Results: The majority of students 53% who responded to the survey are prospective buyers. Fig 7

Current status of the participants

Results: As we previously stated, the proportion of persons without a car was higher since students were more involved than professionals in the workforce and other settings. Fig 8

Type of car variant they prefer when they buy in future

Results: Several factors contribute to why people may prefer petrol cars over electric vehicles (EVs). While the popularity of EVs is growing, there are still certain challenges and considerations that influence consumer choices. Here are some common reasons why individuals may currently prefer petrol cars:

- Infrastructure Concerns
- Price range anxiety
- Initial Cost
- Charging Time
- Longer Refuelling Time
- Limited Model Variety
- Limited Model Options
- Uncertain Resale Value
- Technological ConcernsLess Government Incentives

Preference of price range when participants buy in future:

Results: The preference for mid-range cars over luxury cars among Indian consumers can be attributed to several economic, cultural, and practical factors. Here are some key reasons why many Indians tend to opt for mid-range cars:

- Affordability
- Fuel efficiency
- Average road Conditions
- Efficient resale Value
- Low maintenance Costs
- Demographic preferences
- Limited luxury infrastructure

While these factors contribute to the prevalence of mid-range cars in India, it's essential to note that the luxury car market is growing, and changing economic dynamics and consumer preferences may lead to shifts in purchasing patterns over time.





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Preference of adding AI in cars:

Results: People appreciate and embrace AI (Artificial Intelligence) in their cars for several reasons, as it brings various benefits and enhances the overall driving experience. Here are some key reasons why individuals love AI in their cars: Fig 8

- 1. Enhanced Safety Features
- 2. Driver Assistance and Automation
- 3. Improved Navigation and Traffic Management
- 4. Voice-Activated Controls
- 5. Personalization and Customization

Are the participants willing to spend an additional one lakh rupees to deploy AI?

Results: The data indicates that most people are willing to invest in AI for their automobiles, which suggests that people are prepared to make the rapid transition from conventional to AI-enabled vehicles. Fig 9

Getting deeper into the data

We divided buyers into three categories: LOW, MID, and HIGH. According to our survey, 71% of Indians choose low- and mid-range cars over high- or luxury-class vehicles, indicating that low- and mid-range users dominate the market. However, the turning point came when individuals in all three categories expected their cars to include AI functions, and many of them were prepared to pay for them. Let's examine each category separately. Fig 10

Category -1:

Results: Here we can see that nearly 72% of the people from low budget range expecting AI feature to be installed in their cars and 61% are ready to invest a lakh more for it Fig 11

Category -2:

Results: Here we can see that mid budget people are also interested to have AI features in their car. 58% likes to have it and out of 58% nearly 60% likes to invest a lakh more for AI feature is quite impressive. Fig 12

Category -3:

Results: Over 80% of people who prefers to buy luxury cars needs AI in their cars. Out of 29% who voted as they prefer luxury cars than low are mid range cars ,more than 70% in high budget category are ready to invest more for AI in their cars says that any company which does not integrate AI cannot survive in Indian market Fig 13

CONCLUSION

The automotive industry has undoubtedly witnessed a significant transformation in the design, manufacturing, and user experience of automobiles due to the incorporation of Artificial Intelligence (AI). We have examined the several uses of AI in the automobile industry through our examination of the literature, as well as how it can revolutionize market analysis by utilizing collected data. This article provides a complete understanding of artificial intelligence (AI), how it has been implemented in the automotive industry, and Why we need to integrate AI into low- and midrange cars in a way that benefits both the manufacturer and the customer. The role of the Indian market in the global economy and the most recent car sales data have been analyzed and visualized. AI has ushered in a period of innovation and growth, from improving safety features through Advanced Driver Assistance Systems (ADAS) to enabling autonomous driving capabilities. AI algorithms enable cars to comprehend complicated settings, make wise decisions, and instantly adjust to changing road conditions by utilizing enormous datasets. Moreover, market analysis in the automobile industry has been transformed by the combination of AI and collected data. When it comes to forecasting market trends, understanding customer behavior, and guiding decision-making processes, raw data is a strategic asset. Large-scale datasets powering machine learning algorithms give stakeholders a better understanding of market dynamics and help them formulate competitive strategies. In conclusion, artificial intelligence (AI) is having a significant and wide-ranging impact on the automotive sector. This signals the arrival of a time when intelligent cars will be able to anticipate and adapt to changing societal demands in addition to



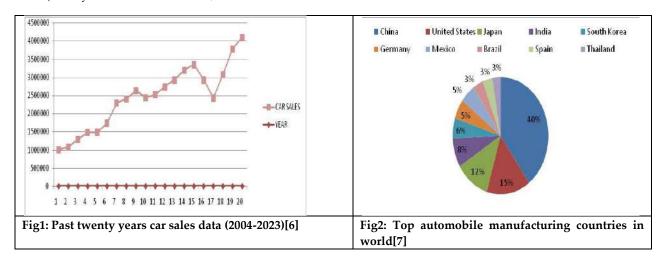


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managing highways. AI will surely continue to play a major role in determining the direction of mobility in the future, guiding the development of a safer, more intelligent, and environmentally friendly automobile ecosystem and by incorporating AI into low- and mid-range cars, any company that makes automobiles, especially cars, may increase their profits and thrive in the Indian market.

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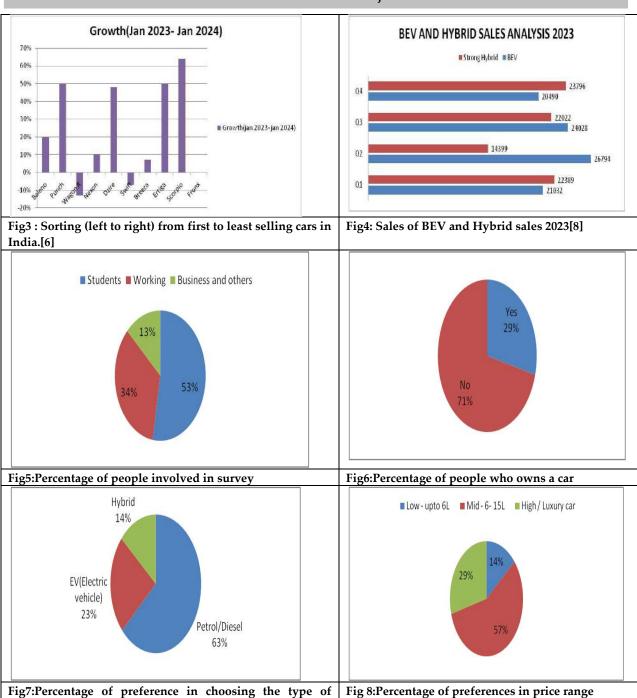


vehicle



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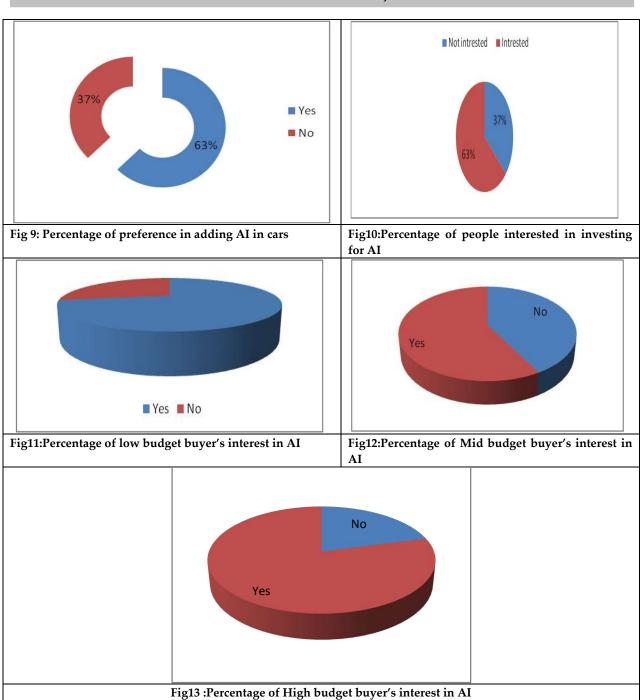
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RESEARCH ARTICLE

Energy Management Strategy for Hybrid Energy Storage - Electric **Vehicles Based on Intelligent Controllers**

Gundu Venu^{1*}, Giriprasad Ambati², Chamakura Krishna Reddy³

¹Assistant Professor, Department of Electrical and Electronics Engineering, Malla Reddy Engineering College (Autonomous), (Affiliated to Jawaharlal Nehru Technological University Hyderabad), Telangana, India.

²Assistant Professor, Department of Electrical and Electronics Engineering, Vallurupalli Nageswar Rao Vignan Jyothi Institute of Engineering and Technology, (Affiliated to Jawaharlal Nehru Technological University), Hyderabad, Telangana, India.

³Assistant Professor, Department of Electrical and Electronics Engineering, BVRIT Hyderabad of Engineering for Women, Hyderabad, Telangana, India.

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*Address for Correspondence Gundu Venu

Assistant Professor,

Department of Electrical and Electronics Engineering,

Malla Reddy Engineering College (Autonomous),

(Affiliated to Jawaharlal Nehru Technological University Hyderabad),

Telangana, India.

Email: gunduvenu83@mrec.ac.in



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ABSTRACT

Developing an Energy Management Strategy (EMS) that takes into account the Electrical Vehicle's (EV's) power distribution between the battery and ultracapacitor helps lessen the EV's power usage and prolong battery life. Therefore, the goal of this paper is to create a Fuzzy Logic Controller-based EMS for EVs that takes battery degradation into account. The hybrid energy storage electric vehicle model is developed first for EMS verification. Battery deterioration modelling trials are complete in the meantime. After that, a Hybrid Energy Storage System's (HESS) power is distributed sensibly using a rule-based control strategy. By comparing the proposed EMS to the existing EMS and the findings, it becomes clear that the former uses less energy and causes less battery degradation. The features of the HESS serve as the basis for the experiment's design, execution, and modelling of battery deterioration. Based on the model, the proposed work tested in MATLAB/Simulink a rule-based EMS and a PMP energy management method that takes battery degradation into account. Considering battery deterioration, the





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limited energy of hybrid energy storage electric vehicles is divided equitably and utilized to maximize energy management.

Keywords: electric vehicle; hybrid energy storage system; energy management strategy; fuzzy logic Controller; battery degradation

INTRODUCTION

The research and development of hybrid energy storage systems, which can increase the efficiency and power of electric vehicles, has been receiving greater attention recently. Both batteries and ultracapacitors play significant roles in a hybrid energy storage system. Lithium-ion battery shortages can be mitigated by using ultracapacitors because of their high-power density, rapid charge and discharge times, and multiple-cycle life [1]. On the other hand, the battery's power and efficiency would gradually degrade over time due to electrochemical processes and the battery's state [2]. In addition, the energy management system in a hybrid electric vehicle is responsible for controlling and balancing the power supplied by the car's power battery and ultracapacitor [3-4]. To maximize vehicle efficiency and prolong battery life, the energy management system must take battery degradation into account while providing the required power. Consequently, the primary difficulty of an energy management approach for hybrid energy storage electric vehicles is figuring out how to fairly divide the power from the power battery and the ultracapacitor. Hotspots in the study of energy management for hybrid energy storage electric cars include the design of energy management strategy, the construction of the battery decay model, and the matching of ultracapacitors and DC/DC converters. Battery ageing factors have been the basis for numerous models published by domestic and international academics depicting the decline in battery life and performance over time. The evaluation models are used to offer energy management techniques that take battery degradation into account [5]. There are now three types of models used to describe the decline in lithium-ion battery capacity: mechanism models, equivalent models, and empirical models. The physical and chemical features of the battery provide the basis for the mechanism model, which aids in the investigation of the causes of battery capacity loss. Mechanism models are typically employed in battery research rather than control problems [6] due to the complexity of the model creation procedure and the difficulty of acquiring the electrochemical parameters included in the model. The correctness of the equivalent circuit model is affected by several external factors, and the electrical components are susceptible to losses.

The model accuracy requirements of this investigation can be met with the use of an empirical model, the establishment and computation of which are, however, rather straightforward [7]. Considering the battery capacity decline and the physical interaction of many parameters, a considerable amount of test data on battery life is employed for formula-fitting construction [8]. Battery state of charge, state of charge, and ultracapacitor state of charge are all examples of inputs. When discharging a battery or capacitor, the power is determined by the battery power distribution coefficient. Although this management strategy's foundation is straight forward to pick up, the control principles behind it necessitate extensive expertise. Complete utilization of composite power supply with pinpoint accuracy is unachievable. An onboard lithium-ion battery system's thermal safety and degradation were highlighted in recent research [9] that offered a universal algorithmic framework integrating a model-based state observer and a deep reinforcement learning-based optimizer. Rapid charging, thermal safety enforcement, increased battery life, and computational tractability are all areas where it has proven itself better. Optimizing energy management in rechargeable hybrid energy storage electric cars was the focus of [10], which also offered a battery model. This model predicts the rate of decline in battery health as the state of charge (SOC) and pack temperature are changed. The numerical multi-objective optimum control problem is solved by using SDP and PSO to strike a balance between power consumption and battery life while meeting the needs of the vehicle. Efficiencies in battery life and power consumption were analyzed.





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Hybrid Energy Storage Electric Vehicle Modelling

In this paper, a closer look at the parallel hybrid energy storage system (HESS) is shown in Figure 1 which consists of a controller, motor, and ultracapacitor in addition to a battery and a DC/DC converter.

Vehicle Modelling

In order to simulate a hybrid electric car, it is necessary to analyze a simulation experiment involving energy management. That's why this research constructs a model car, complete with a driver model, a vehicle longitudinal dynamic model, a motor model, and a HESS model. While the other models are constructed using mathematical modelling, the motor model is generated by utilizing simulation results. In this paper, the UC module is connected to the DC bus via a bidirectional DC/DC converter, while the battery module is connected directly to the DC bus, resulting in a UC semi-active hybrid topology. The DC/DC converter can regulate the UC's output power to deliver the maximum amount of energy to the vehicle while keeping the system's overall power consumption to a minimum.

While the other models are constructed using mathematical modelling, the motor model is generated by utilizing experimental methods. In this research work, the UC module is connected to the DC bus via a bidirectional DC/DC converter, while the battery module is connected directly to the DC bus, resulting in a UC semi-active hybrid topology. Also, the DC/DC converter may regulate the UC's output power to deliver the vehicle's maximum power while minimizing the system's overall power draw and extending the battery's service life. The HESS model is calibrated using the battery, UC, and DC/DC factors and characteristics. It is the Rint equivalent circuit that is chosen by the battery model. The model's power battery has an open circuit voltage and internal resistance that vary with state-of-charge (SOC) and temperature, and whose values can be determined with experimentation. Power in the HESS is managed by a buck-boost bidirectional DC/DC converter, and a model of this converter is developed based on its operating principle and specifications.

Energy Management Strategy Based on Fuzzy Control

In energy management strategy, the primary function is to regulate power distribution in hybrid electric vehicles and all other electric vehicles that are operated with battery management. As a rule-based control approach, fuzzy control is very flexible and reliable. An important problem in developing energy management strategies is figuring out how to maintain the battery operating conditions in its optimal discharge state. Distribution of the discharge current from the battery and ultracapacitor will accomplish this purpose. It is possible to test the efficacy of the approach by plugging the multiplier for dispersed current into a battery decay model and determining the resulting battery decay rate. In this paper, the notion of the power distribution coefficient K symbolizes the importance placed on battery power, and picking the vehicle demand power, the ultracapacitor SOCuc, and the battery SOCb to achieve the best possible power distribution ratio. This allows for an examination of the fuzzy controller's power distribution coefficient.

$$K = \frac{P_b}{P_{req}} \tag{1}$$

$$P_b = P_{reg}(1-k) \tag{2}$$

where battery power (Pb), UC power (Pc), and vehicle power demand (Preq) are all inputs to the equation.

Based on the hybrid energy storage electric vehicle's operational characteristics, the following control ideas can be derived: when the required power is positive, the hybrid energy storage system outputs power; on the one hand, the ultracapacitor is used to protect the battery from discharging smoothly; on the other hand, it provides instantaneous power to the drive motor. Hybrid energy storage systems can recover energy whenever the power demand is negative. The ultracapacitor restores energy lost as a result of braking, and it also shields the battery from the destructive effects of high-current shocks. Power levels and amplitudes, as well as the emphasis placed on regulation, are distinct between the two modes of operation. It is necessary to design separate control methods for the two modes of operation of the composite power supply. Using the control concepts, it may categorize the control approach into two modes: driving and braking. Fuzzy controller 1 fairly divides the power during the discharge phase, and fuzzy controller 2 determines the charge phase power distribution.





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Control Method

FLC evaluates input data in terms of logical variables with continuous values ranging from zero to one. The operation of the controller in a fuzzy logic control system is based on fuzzy rules generated using fuzzy set theory. Figure 3block diagram of Fuzzy logic controller. The fuzzification, defuzzification, input, and output variables make up FLC. The fuzzy sets are formed using seven membership functions for the inputs error E, change in error CE and the output (Ploss) are shown in Figures5 and 6 and Figure 4 depicts the output.

Simulation Results

Figure 5 depicts the ideal power distribution of the HESS with two distinct energy management strategies under UDDS operating conditions, demonstrating that both strategies can efficiently divide the power of batteries and the ultracapacitor. According to the findings, the PMP energy optimization management technique outperforms the fuzzy control strategy in terms of controlling battery degradation. The optimization method reduces the battery's peak current by 29.5 A, the cell deterioration rate after 300 cycles by 2.33 percent, and the energy consumption rate by 11.72-kilowatt hours per 100kilometers. To sum up, hybrid electric vehicles that employ a PMP-based energy management system that takes battery deterioration into account can significantly postpone battery decline and energy usage, save energy, and protect batteries.

CONCLUSIONS

The purpose of this work is to present an FLC-based energy management approach for hybrid electric vehicles that takes battery degradation into account, with the end objective of assessing the effect of battery decline on the power and economic benefits of electric vehicles. The hybrid electric vehicle model was first established to ensure the strategy's success. To prevent the battery from degrading too quickly and keep the vehicle's power performance stable, the fuzzy control method is utilized to intelligently divide the load between the ultracapacitor and the battery. Lastly, the proposed control method is verified using simulation.

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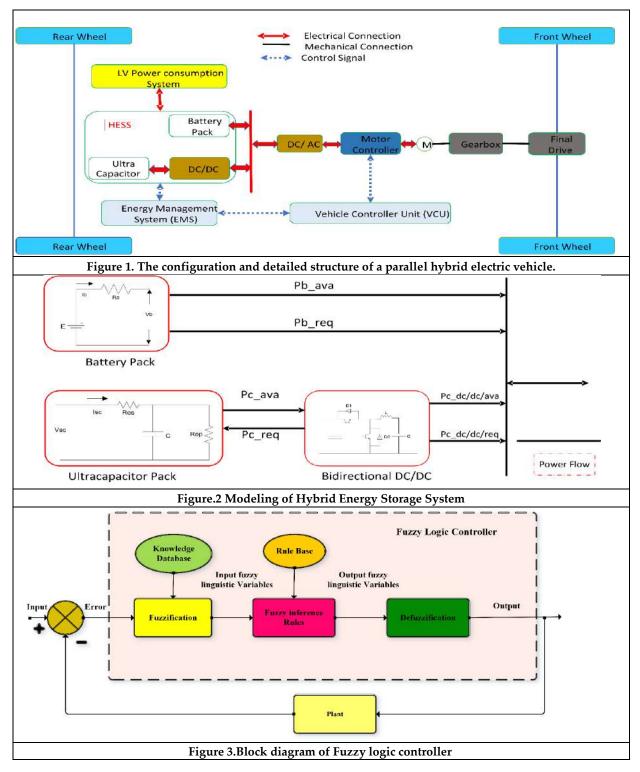
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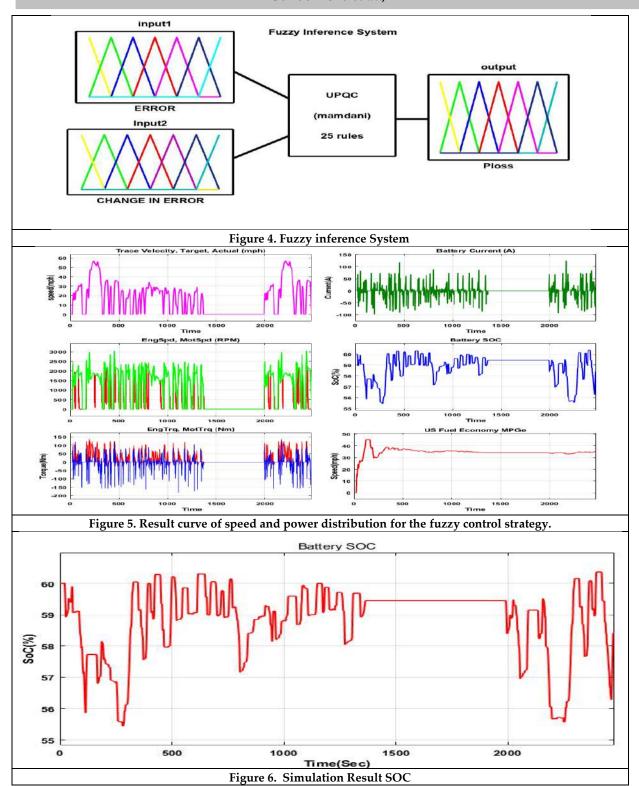
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REVIEW ARTICLE

Current Evidences of Physiotherapy Interventions for Unilateral Neglect in Stroke Individuals: A Systematic Review

Nirmitee Honap¹ and Suraj B. Kanase²

¹Department of Physiotherapy, Krishna Vishwa Vidyapeeth (Affiliated to Deemed to be University) Karad, Maharashtra, India

²Professor and HoD, Department of Neuro Physiotherapy, Krishna Vishwa Vidyapeeth (Affiliated to Deemed to be University) Karad, Maharashtra, India

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*Address for Correspondence

Suraj B. Kanase

Department of Neuro Physiotherapy, Krishna Vishwa Vidyapeeth (Affiliated to Deemed to be University) Karad, Maharashtra, India.

Email: drsurajkanase7@gmail.com

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ABSTRACT

Neglect alone is the inability to respond to external stimuli, including visual, somatosensory, auditory, and kinesthetic sources. The incidence of stroke in the right hemisphere varies between 10% and 82%, and in the left hemisphere between 15% and 65%. Neglect is usually caused by a major stroke in the area of the middle cerebral artery, and the result of neglect will be more severe and persistent pain following damage to the right hemisphere. Different treatments and tools have been developed to measure and solve neglect. Recent data suggest that rehabilitation can be divided into two types of behavior: protecting the hemiplegic limb to reduce ipsilateral site preference or increasing awareness of the source of conflict to support the case. In particular, we recently performed a systematic review of the use of different physiotherapy modalities in the USN complex, including prismatic adaptation, optomotor stimulation, neck vibration, galvanic vestibular stimulation, non invasive brain stimulation, robotic learning, and smooth tracking exercise. The aim of this review is to determine current physical therapy options for neglect in stroke patients. Articles were selected according to main subject headings and analyzed. Clinical studies have provided evidence that physical therapy is effective in improving symptoms of self-neglect in stroke patients.

Keywords: Unilateral Neglect, Stroke, Physiotherapy





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INTRODUCTION

Stroke is the predominant cause of long-term disability worldwide. Consequences of stroke include motor and cognitive impairment, and survivors often develop impairments in activities of daily living (BADL) and independence in activities of daily living (IADL), life satisfaction, and depression. USN is a lack of response to contralateral stimulation that occurs in 25% to 48% of stroke patients. The ability of USN has been studied extensively and its presence has been shown to be associated with less BADL and IADL, fewer threats, longer hospital stays, and better outcomes. Activities to improve the UN are carefully designed, implemented and researched. [1] It is characterized by the inability to express or respond to stimuli from various sources such as visual, somatosensory, auditory and kinesthetic sources. The incidence of stroke is reported to be 10-82% in the right hemisphere and 15-65% in the left hemisphere. A variety of treatments and tools have been developed to evaluate and treat UN problems. Recent data suggest that rehabilitation can be divided into two types of actions: preservation of the hemiplegic limb to reduce ipsilateral preference or increasing awareness of the location of the problem to support the case. It is also difficult to declare which method is the best treatment recommendation because there is no evidence and, interestingly, doctors rarely use scientific evidence.[2]

Unilateral neglect (UN) is a medical term that refers to problems resulting from the failure to report, respond to, or refer to new or useful stimuli occurring on the side of the brain opposite the injury. UN affects 23.5% to 67.8% of stroke patients. However, the most commonly used tests (subtraction task and National Institutes of Health Stroke Scale (NIHSS) item 11) show greater than 30%. The United Nations is an initiative that addresses problems such as long-term homelessness, increased mobility, and decreased family responsibilities, which occur in 30-60% of the population 1 year after a stroke. Although there are many shortcomings in the work of the United Nations, a consensus has not been reached on effective and efficient strategies for measuring and solving this problem. The lack of valid research and effective treatment strategies illustrates the challenge of BM. The UN usually deals with multiple situations rather than the outcome of a single situation. Rather, it is a result of disruptions in the functional connectivity of brain regions associated with auditory, sensorimotor, and visual functions, especially in the frontoparietal network. Therefore, the symptoms are diverse and can occur in different areas (visual, auditory, and sensorimotor) and locations (particularly inside and outside the body).[3] Neglect is often the result of a serious middle cerebral artery stroke, left spatial neglect is more severe, and after a left semicircular stroke, left spatial neglect is more severe, and persistent than on the right.[5]

Spatial neglect has been studied extensively since the 1970s. The first study used a top-down approach, adding unconscious events to non-conscious events. However, the results of these original studies were mixed, which led to the development of alternative methods.[6] Recently, the United Nations has been interested in eliminating consciousness-based neuropsychological problems in space, but its impact has not yet been investigated. Although there is no consensus, active research has demonstrated the effectiveness of prismatic adaptation (PA), transcranial magnetic stimulation (TMS), and transcranial direct current stimulation (tDCS) interventions in the treatment of USN. Additionally, preliminary evaluation demonstrates the potential of electrical vestibular stimulation (GVS) and virtual reality (VR) methods in USN rehabilitation. Although many studies on SSN were published in the 20th and early 20th centuries, neuropsychological research has decreased in the last 10-15 years. The first reason could be twofold. Advances in neuroimaging and brain stimulation have recently led to an increase in the number of neurologists using these techniques. A new tool for studying the neural basis of spatial cognition in the healthy brain. Second, pharmacotherapy improves clinical and neuropsychological outcomes in stroke patients.[7] Many different methods have been used for a long time to improve the symptoms of neglect.

Many studies have been published on each method. Since this is not a medical negligence review, we will focus on the best treatments mentioned, although there are other good but less effective methods. Learn to treat this problem like eye patching,[8] caloric vestibular stimulation,[9]. Visuomotor imaging,[10] mirror therapy,[11] TENS, [12] photokinetic stimulation [13 – 15] and Constraint-induced movement therapy. [15 – 17] Specifically, we reviewed





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research on the use of prism adaptation, noninvasive brain stimulation, and virtual reality in USN rehabilitation. Therefore, we use the words "neglect", "treatment", "prism adaptation", "tDCS", "vestibular stimulation", "TMS", "TBS" and "virtual reality" together.

METHODOLOGY

Search Strategy

We performed A systematic review for current physiotherapy interventions for unilateral neglect in stroke patients. For the purpose of this review, an electronic search for relevant articles was done using Google Scholar, PUBMED, MEDLINE, Pedro, Research Gate and CINHAL database from to 2013 to 2023 was done where in MeSH search terms such as cerebrovascular accident OR stroke; neglect; visuo-spatial neglect; visual neglect; unilateral neglect; and hemisphere neglect free words were used. The search was bounded to RCTs involving adults aged 19 or over. We included all RCTs that sought to identify the effectiveness of any type of physiotherapy intervention in Unilateral neglect in adult stroke patients diagnosed by clinical examination and/or classical neuropsychological tests. Only studies which reported only physiotherapy intervention were included. Observational studies and case reports, systematic reviews or meta-analyses, and cross-sectional studies were excluded. Additionally, searches for relevant books were performed manually. Articles were selected based on experience, self-awareness, and reflective practice.

Study selection

Review selection criteria include

- 1. Physiotherapy intervention for patients suffering from unilateral neglected stroke.
- 2. Effectiveness of physical therapy for unilateral neglect in stroke.
- 3. Research on stroke in adults.

Data extract

All stages of study selection were assessed against inclusion criteria. Study titles and abstracts were screened by reviewers. Relevant articles were reviewed in full text and included if they met the inclusion criteria. The following data were extracted from relevant articles, including purpose, study design, study population, study period, physiotherapy intervention, selected outcome measures, and main outcomes. Literature review (Table 1)

RESULT

The search identified 10 relevant studies. Inclusion criteria for further analysis were met. A new treatment method combining neck muscle vibration and transcranial stimulation appears directly in the routine treatment of patients with left hemi-neglect. HEMISTIM protocol of randomized trials led by Sarah Millot, Jean Marie et al., 2023. The purpose of this study was to evaluate immediate and long-term improvements through a novel relationship between left cervical muscle oscillation (NMV) and bipolar transcranial. Direct current stimulation (tDCS) of the posterior parietal cortex during occupational therapy in patients. He left USN. middle. Participants were divided into four groups: control, left NMV, sham left NMV-tDCS, or anodal tDCS left NMV. NMV and tDCS were used for the first 15 minutes of treatment, 3 days a week for 3 weeks. USN was assessed at baseline, at the end of the first study, after the first and third weeks of policy, and 3 weeks after policy end. The primary outcome of this study was change in functional scores on the Katherine Bergo Scale. Secondary outcomes included five tests examining USN neuropsychological variables. The vibrating stimulator is placed on the skin over the belly of each trapezius muscle and secured with a belt. The results of this multimodal approach also provide interesting insights into the hemispheric perspective and demonstrate the importance of right hemisphere or other "hemispheric" support. [18] Thai Regina, Helio Rubens and others. the 2022 study, "Non-invasive brain stimulation reduces unilateral spatial neglect after stroke," examines the impact of physical therapy following anodal and cathodal transcranial direct current stimulation (A-tDCS and C-tDCS, respectively) on stroke recovery. Posterior USN This double-blind, randomized pilot study was conducted in patients with OSI after ischemic stroke. Randomization was stratified by





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Behavioral Inventory-Inventory (BIT-C) and Catherine Belgo Scale (CBS). Twenty minutes of tDCS was followed by topical treatment for 7.5 weeks. The biggest advantage is the USN degree evaluated by BIT-C. Secondary outcomes were various CBS scores, stroke (National Institutes of Health Stroke Scale [NIHSS]), disability (modified Rankin Scale), autonomy (Barthel Index, Functional Independence Scale), and quality of life (five-dimensional EuroQol). personality survey). For the primary outcome, pairwise post hoc comparisons were performed using Bonferroni correction. The study results showed greater improvement in BIT-C scores after A-tDCS intervention compared to sham (mean difference [MD] = 18.4). There were no significant differences between groups in secondary outcomes. However, no improvements were observed in activities of daily living, stroke, dependency, independence, functional independence, or quality of life. Both stimulation methods have been shown to be safe. In conclusion, A-tDCS was shown to have a significant effect on physical therapy and reduce USN levels in the subacute phase of stroke. [19] Dr. Hyunse Choi and Dr. Bomin Lee conducted a study titled "Comprehensive Integrated Prism Adaptation and Neck Vibration Intervention for Unilateral Neglect in Chronic Stroke Patients" in 2022 to study the effectiveness of an integrated prism adaptation (PA) rehabilitation program. target.) and neck vibration (NV) due to unilateral neglect in stroke patients. 36 patients were divided into the PA NV group (group A, n=12), NV only group (group B, n=12), and PA only group (group C, n=12). The intervention was conducted for 50 minutes a day, 5 times a week, for 4 weeks. Traditional activities include joint mobility, functional training, and ADL training. Joint exercises include passive range of motion (ROM), assisted active range of motion, and resistance training.

Vocational training includes many activities such as assembling building blocks and assembling rings, as well as the next steps, including the role of the patient. ADL training includes eating, dressing, and using the bathroom. PA uses triangular glasses with a left-angled base that provides a 10° deviation of the visual axis to the right, so the user sees the object as if it has shifted 10° to the right. Vibratory stimulation is used for the neck extensor muscles. Vibratory stimulation is applied to the upper sternomastoid and osteomastoid muscles, located approximately 6 cm from the cervical spine. Allow the patient to rest for 30 seconds between each 2-minute dose increase. The Albert and Catherine Bergo scale was used to assess the impact of each intervention on personal neglect, and the modified Barthel index was used to assess the impact of ADL. After the intervention, all three groups experienced a decrease in neglect and improvements in activities of daily living (p < 0.05). Importantly, the reduction of neglect alone in group A (PA NV) was higher than in the other groups (p < 0.05); However, the improvement in ADL was not different between the three groups (p>0.05). This novel intervention involving PA NV is recommended for the treatment of chronic stroke patients with clinical failure. [20] Effects of mirror visual feedback on neglect in poststroke patients: Kenneth Fong, K.H.Ting, et al (2022) conducted a preliminary randomized controlled trial in which a total of 21 subacute patients with left spatial neglect after right hemisphere stroke received a randomized clinical trial. For 3 groups: MVF, Sham 1 (view of the hemiplegic arm through a clear glass with both arms in motion) and Sham 2 (using a cover mirror). The 3-week treatment for each group included 12 physical exercises on the hemilegic arm, graded according to the severity of the injured arm.

A blind test was conducted before/after surgery and 3 weeks later. The results show that MVF does not have a significant effect on Simulation. However, as indicated by the diagonal line, MVF was significantly more effective than Sham 2 (p = 0.022). In contrast detection, MVF was slightly better than using a protective mirror at distinguishing numbers on the left and right sides of the page (p = 0.013; p = 0.010), indicating that MVF had little effect on reducing allocentricity. This study confirmed the superiority of MVF over using coverslips as a method to reduce spatial sloppiness and allocentric symptoms, but the results were not obtained. An alarm was seen when compared to two arms moving inside the transparent glass.[21] AlessioFacchin, GuisiFigliano and colleagues conducted a comparative study of prism-adapted optokinetic stimulation in unilateral spatial neglect rehabilitation in 2021. The aim of this study was to compare the effectiveness of two methods in an underserved patient group. Transitional design. PA and OCS were administered to 13 posttraumatic brain injury patients with unilateral spatial neglect. Each treatment was administered 10 times twice daily, with each patient receiving 2 treatments in a alternating sequence (e.g. PA followed by ACS or vice versa). Neuropsychological testing; It was conducted before the first session (T1), at the end of the first session/start of the second session (T2), at the end of the second session (T3) and 2 weeks after the end of treatment. (T4). Regardless of the type of training, both methods provide significant





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improvements in T2 diagnosis. The results showed that PA or OCS had a significant effect on neglect in patients with right hemisphere brain injury, mainly in the first phase of treatment. There is no difference in treatment, so it can be used in treatment according to the patient's needs. [22] Effect of robotic training on the left side of hemi-spatial neglect in adults with stroke: A controlled study conducted in 2021 by Pilot and Park JH aimed to evaluate the effect of robot-assisted training on hemi-spatial neglect in adults with stroke. Participants were assigned to an experimental group (EG) (n = 12) receiving left hand robotics training or a control group (CG) (n = 12) receiving symptom care treatment. Each participant completed 20 sessions over 4 weeks. The Line Division Test (LBT), Albert Test, and Catherine Bergo Scale (CBS) were used to examine the impact of unilateral spatial neglect. Results were analyzed before and after 20 training sessions. After intervention, LBT showed improvement in the Albert test and CBS EG, while LBT and CBS showed improvement in the CG but did not improve in the Albert test. These results indicate that robotic training is effective in improving unilateral spatial dysfunction in adult stroke patients compared to non-stroke patients. The study concluded that the use of robotic limbs could help improve unilateral spatial neglect. [23] Efficacy of cervical taping in the treatment of unilateral spatial neglect in stroke patients: a pilot single-blind randomized controlled trial conducted in 2019 by Valentina Varalta and Daniele Munari.

The purpose of this study is to evaluate the effectiveness of therapeutic cervical taping in stroke patients. Visual-spatial ability, neck movement, and kinesthetic sensitivity in chronic stroke patients with hemispatial neglect. After randomization, 12 chronic stroke patients with hemispatial neglect underwent real (treatment group) or sham (control group) neckbanding for 30 consecutive days. The results are as follows: star removal test; neck range of motion; probation denial letter; Comb and razor control; Cervical spine joint dysfunction before and after one-month closure. Comparison between groups revealed a significant difference in neck joint error only after treatment (p = 0.009). Preliminary results support the hypothesis that neck taping may increase cervical kinesthetic sensitivity in chronic stroke patients with hemispatial neglect. Additional studies are needed to confirm our results and investigate the beneficial effects of therapeutic elastic tape on visuospatial abilities in stroke patients with hemispatial dysfunction[24] Kim SB, Lee KW, and Lee JH conducted a study on "The Effect of Combination Therapy of Robots and Low-Frequency Repetitive Transcranial Magnetic Stimulation on Hemispatial Neglect in stroke Patients" in 2018 to investigate the treatment of upper extremity pain by connecting robots together. Effects of cranial magnetic stimulation (rTMS) on unilateral spatial neglect in stroke patients. Patients with hemispatial neglect after right hemisphere stroke were divided into rTMS only group, robot only group and combined group.

Each group received a well-structured neglect treatment plus additional treatment for each group. The robotic group received robotic therapy, and the combined group received both treatments. The treatment effect was evaluated using the Visual Perception Test-3 (MVPT-3), Line Bisection Test, Asterisk Elimination Test, Katherine Bergo Scale (CBS), Korean version of Mini-Mental SE (MMSE), and Basel Index (K-). MBI). These parameters were measured before and after treatment. Ten patients were selected in each group. There were no remarkable differences in values between the three groups. After two weeks of treatment, MVPT-3, Linear Test, Star Elimination Test, CBS, MMSE, and K-MBI scores significantly improved in all groups. However, there was no significant difference in the rate of change between groups. It can be concluded that the combined effect of robotic therapy and low-frequency rTMS therapy in the treatment of hemispheric neglect does not differ depending on the treatment method. The results of this study showed no difference. Additional studies with larger patient groups are needed to evaluate the effectiveness of these treatments. [25] Volkening K, Kerkhoff G, Effects of repetitive galvanic vestibular stimulation on spatial neglect and vertical perception - a randomized controlled trial. They conducted the smallest randomized controlled trial. Twenty-four patients received 10 to 12 sessions per day, 5 days per week. The CL- and CR-GVS groups received 1.5 mA current stimulation for 20 min, whereas the Sham-GVS group received CL-GVS for only 30 s. Simultaneously, all patients were given clinical examples of visuomotor tasks (VMT) and visual training (VST). Test outcomes (ignorance test, visuo-tactile search, visual and tactile vertical) were assessed before the intervention, immediately after the intervention, and 2 and 4 weeks later. Results showed that neither our standard treatment nor the combination of standard treatment and GVS improved neglect symptoms. ²⁶ Appropriate eye movement training promotes recovery from hearing and vision loss: a randomized controlled trial by Georg Kerkhoff and Stefan Reinhart Objective: To compare the effectiveness of CBT and visual stimulation (VST) on hearing and vision loss in





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stroke patients. Fifty patients with left-sided hearing and visual impairment were excluded from the study. Twenty-four patients received CBT and 21 patients received VST. Five patients (4 VST, 1 SPT) were absent. Each group received one hour of neglect treatment for a total of five hours over five consecutive days. Outcome measures for visual impairment (digital clipping, visuomotor separation, sentence reading) and hearing impairment (intermediate vision) were assessed twice: before treatment, after treatment, and two weeks later. The SPT group showed good eye movements when following left-hand stimuli. The VST group systematically scanned identical but static stimuli. Both groups were divided into subgroups to study the effects of mild and severe neglect. Results: There were no differences between the two groups in pre-treatment demographic variables or level of neglect (hearing/visual). After treatment, the CBT group showed, on average, significant and sustained improvements in all visual and auditory measures. No significant changes in vision or hearing loss were noted after VST. Additionally, treatment effect sizes following PCT (Cohen's d) for visual and hearing impairment were greater than effect sizes following VST treatment, regardless of mild and severe neglect. The study concluded that contralateral smooth tracking training was more effective than combined treatment, resulting in mild and major neglect. [27]

DISCUSSION

This review summarizes 10 years of current data on physical therapy for neglect alone in stroke patients published in the published literature in clinical trials. Randomized controlled trials provide most of the evidence in this study. However, there are some differences between the recommendations and the research results. These differences may be due in part to the methods used in the studies reviewed, the types of research selected, the different characteristics of the studies examined, and/or the brand target or purpose of these studies. Therefore, there is a need for quality studies to evaluate the effectiveness of physical therapy alone in low-income stroke patients. The small number of articles reviewed may represent a limitation of this review. Finding other areas of conflict in stroke patients will expand our analysis and will not influence future recommendations and recommendations for physical therapy. However, Google Scholar and Pubmed were chosen because they are informative and support the purpose of the current review, which is to evaluate clinical trials for the treatment of this disease. Sarah Millot, Jean Marieet conducted a randomized controlled trial. et al combined neck muscle stimulation and transcranial direct current stimulation with respiratory rehabilitation in patients with left unilateral neglect - the HEMISTIM protocol. Findings suggest that using this modification in the treatment process may improve outcomes with rehabilitation. In fact, noninvasive brain stimulation is known to modulate cortical excitability and is therefore an ideal tool to promote longterm neuroplasticity. The results of the study "Non-invasive brain stimulation reduces unilateral spatial neglect after stroke", conducted by Tais Regina, Helio Rubens, showed that A-tDCS has a positive effect on the treated body, thereby reducing the risk of stroke USN19. in the subacute stage.

Hyun-Se Choi and Bo-Min Lee conducted a study on the effects of a combination of prism adaptation and neck vibration in the treatment of unilateral neglect in stroke cases. PA NV is also among its benefits. ²⁰. Effects of Specular Visual Feedback on Spatial Neglect After Stroke KH Ting, a preliminary randomized trial conducted by Kenneth Fong, concluded that optimal visual feedback using glass covers as a recording language reduced neglect and reduced allocentric symptoms. Prismatic adaptation and optokinetic stimulation can be used in treatment on a case-by-case basis²¹. The results of the study conducted by AlessioFacchin, GuisiFigliano and colleagues in 2021 on the comparison of the transition from prisms to photodynamic excitation in the improvement of negativities in the blind spot well, revealed that PA or OKS believe that negligence. It was about events. It is certain that he is in the first stage of treatment for right brain damage. Because no difference was found between treatments[22]. Recently, left-handed robotic training affected hemispatial neglect in paralyzed adults. Park JH conducted a pilot and randomized controlled study in 2021. Results showed that EG had a small benefit compared to CG in all aspects of development; Therefore, robotic training may help improve the treatment of hemispatial neglect in adults with habitual neglect. Compared to conventional treatments for stroke. This study concluded that robotic support may help improve hemispatial neglect [23]. Effect of Neck Taping in the Treatment of Hemispatial Neglect in Patients with Habitual Stroke Valentina Varalta conducted a randomized controlled trial with Daniele, Munari. The main findings support the





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argument that neck taping may increase cervicofacial kinesthetic sensitivity in stroke cases with hemispatial neglect. This study also noted that further research is needed to strengthen the results and investigate the positive effects of elastic band therapy on visuospatial ability in stroke with hemispatial neglect[24]. In 2018, SB Kim conducted a study on the effect of combined robotic therapy and low-frequency repetitive transcranial charismatic stimulation on hemi-spatial neglect in stroke patients and concluded that the combination of robotic therapy and low-frequency rTMS combination of hemi-spatial neglect. The treatment effect is no different from the effect of treatment alone. The results of this study do not prove the superiority of any of the three treatments. More studies involving more patients are needed to evaluate the effectiveness of these treatments. [25] Volkening K, Kerkhoff G, Keller I conducted a randomized sham-controlled study on the effects of repeated vestibular stream stimulation on spatial neglect and vertical perception. The results showed that neither standard treatment nor the combination of standard treatment and GVS significantly improved neglect symptoms. [26] Training to improve eye function promotes recovery from auditory and visual neglect The study was conducted by Georg Kerkhoff and Stefan Reinhart. This study concluded that contralateral, uniform learning would produce superior multimodal treatment in minor and major neglect conditions. [27]

CONCLUSION

The articles that were previously examined gave a summary of the most recent research on physiotherapy interventions for unilateral neglect in post-stroke patients. The evaluated studies provided evidence that the described physiotherapy techniques are useful in providing significant benefit to patients with unilaterally neglected stroke. If physical therapy includes neck muscle stimulation, direct transcranial stimulation, prismatic adaptation, optomotor stimulation, mirror visual feedback, repetitive vestibular galvanic stimulation, and robotic training, it may be useful to minimize neglect of the hemiplegic side.

CONFLICT OF INTEREST: None

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Table no. 1 Summary of review of literature

Sr	Title of study	Aim of study	Study design	Sample	Study	Outcome	Intervention
1	Innovative therapy combining neck muscle vibration and transcranial direct current stimulation in association with conventional rehabilitation in left unilateral neglect patients. Sarah millot,Jean Marie et.al(2023)	To assess immediate and long term recovery induced by innovative association of left side neck muscle vibration and anodal transcranial direct current stimulation in patients with left unilateral spatial neglect	Randomized clinical trial	size	3 days a week for 3 weeks . Total 45 min of therapy	Catherine bergego scale Fluff test Gainottis drawing test Map of France test	Neck muscle stimulation using a vibrator placed over sternocleidomastoud followed by transcranial direct current stimulation
2	Non invasive brain stimulation can reduce unilateral spatial neglect after stroke. Taisregina,Heli orubenset.al(20 22)	To examine the effects of physical therapy after anodal and cathodal transcranial direct current stimulation to improve visuospatial and functional impairment in individuals with unilateral neglect after stroke.	Randomized control trial	n=51 anodic group= 15 cathodi c group= 15 sham grp=16	Total duration=7 .5 weeks Total sessions =15 given 2 times per week	BIT-C – Behaviou r inattentio n test- conventio nal CBS- Catherine bergego scale	A direct current was delivered by a battery-powered device using 2 pairs of surface saline-soaked sponge electrodes (5cm 5cm). Each side of the sponge was soaked in saline solution. The study mode was designed to encode the sham and active stimulation attempts using 5-digit codes





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3	A complex intervention integrating prism adaptation and neck vibration for unilateral neglect in patients of chronic stroke. Hyun se Choi,Bo min lee (2022)	To investigate the effects of complex rehabilitative programme that integrates prism adaptation and neck vibration for unilateral neglect in patients of chronic stroke.	Single blind Randomized control trial	n=36 PA+NV = 12 Only NV=12 Only PA=12	50 Min per day 5 sessions per week for 4 weeks.	Alberts test Catherine Bergego test Barthel index	Prism adaptation using triangular prism goggles was given. In NV,vibratory stimuli was applied to neck extensors using a vibration stimulator.The patient was given 30s rest after every 2 min performance during the 20 min session.
4	The effect of mirror visual feedback on spatial neglect for patients after stroke: A randomized controlled trial. Kenneth Fong,K.H.Tinge t.al (2022)	To investigate the effects of mirror visual feedback on reduction of spatial neglect for patients with stroke	Randomized controlled trial	n=21	3 week treatment. 12 sessions-4 per week for 3 weeks each lasting for 30 min	Behavior al inattentio n test. Gap detection test. Catherine Bergego scale	In group 1,mirror was used and patient was asked to imitate the movements with the affected limb. In group 2, the mirror was covered by cloth repeating same procedure.
5	Prism adaptation to optokinetic stimulation comparison in rehabilitation of unilateral spatial neglect. AlessioFacchin, GuisiFiglianoet .al (2021)	To compare the effectiveness of these two methods in a group of unilateral neglect patients.	Randomized controlled trial	n= 12 prism adaptat ion=6 optokin etic stimula tion = 6	Each treatment was applied for 10 sessions – twice a day for 5 days a week	Letter cancellati on test Star cancellati on test Copy drawing test Sentence reading test	Prism adaptation was performed first followed by optokinetic stimulation. In the second group , treatment was reversed.
6	The effects of robot assisted left hand training on hemispatial neglect in older patients with chronic stroke: A pilot and Randomized control trial. Park JH (2021)	To investigate the effects of robot-assisted hand training on hemispatial neglect of older patients with chronic stroke.	Randomized control trial.	n=24 12=exp eriment al group 12=cont rol group	20 sessions 5 times a week for 4 weeks. Each session for 30 min	Line bisection test. Albert test. Catherine Bergego scale.	Robotic assisted training was given to left hand using AmadeoRobot.the motion was assisted by robot and the individuals level of function.



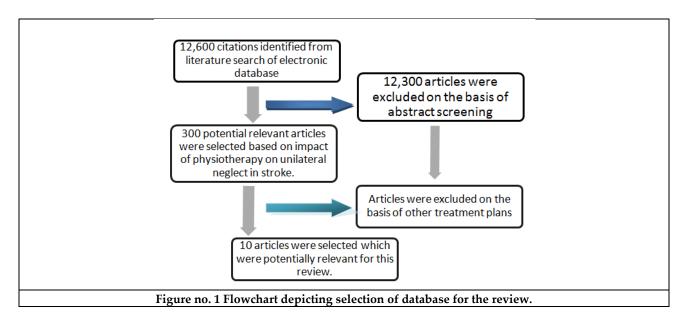


7	Effect of neck taping in the treatment of hemispatial neglect in chronic stroke patients:A pilot single blind Randomized control trial. Valentina Varalta, Daniele Munari (2019)	To assess the effect of therapeutic neck taping on visuospatial abilities, neck motion and kinesthetic sensibility in chronic stroke patients with hemispatial neglect.	Single blind Randomized control trial	n=12 6=treat ment group 6=contr ol group	30 days Tape was replaced every 4 days	Star cancellati on test. Letter cancellati on test. Comb and razor test.	The tape strip was applied from mastoid bone to clavicle with sternocleidomastoid kept in position of maximum stretching-15-25% stretch.
8	Effect of combined therapy of robot and low frequency repetitive transcranial magnetic stimulation on hemispatial neglect in stroke patients. SB Kim, KW Lee (2018)	To investigate the effect of upper limb rehabilitation combining robot with low frequency repetitive transcranial magnetic stimulation(Rt ms)on unilateral spatial neglect in stroke patients.	Single Blind Randomized control trial	n=30 10=Rtm s only group 10=rob ot	30 min a day 5 days a week for 2 weeks.	Motor free visual perceptio n test. Line bisection test. Star cancellati on test	
9	Effects of repetitive galvanic vestibular stimulation on spatial neglect and verticality perception-a randomized controlled trial. Katherina Volke ning, Georg Kerkhoffet.al (2016)	To compare the effects of repeated galvanic stimulation on spatial neglect and verticality perception.	Double blind Randomised controlled trial	n=24	10-12 sessions daily 5 days per week	Neglect test Visuo tactile search test	Participants received galvanic stimulation and simultaneously received a standard therapy of smooth pursuit eye movement training followed by visual scanning training.
10	Smooth pursuit eye movements training promotes recovery from	To compare the effects of SPT and visual scanning therapy on	Randomised controlled trial	n= 24 for SPT n=21 for VST	Each group received 1 hr session for 5	Digit Cancellat ion Visuoper ceptual	The SPT group practiced smooth pursuit eye movements while trackinf stimuli moving leftward.





auditory and	auditory and		consecutiv	and	The VST group
visual neglect:	visual neglect		e days for	motor	systematically scanned
A randomized	in chronic		3 weeks.	line	the same but static
controlled	stroke patients			bisection	stimuli.
study.	with neglect.			Paragrap	
Georg				h reading	
Kerkhoff,					
Stefan Reinhart					
et.al (2013)					







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RESEARCH ARTICLE

Degree-Based Degree-Splitting and **Evaluation** Salbutamol's of **Topological Indices**

R.Revathy*

Assistant Professor, Department of Mathematics, Sri S.Ramasamy Naidu Memorial College, Sattur (Affiliated to Madurai Kamaraj University) Madurai, Tamil Nadu, India

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*Address for Correspondence R.Revathy

Assistant Professor, Department of Mathematics, Sri S.Ramasamy Naidu Memorial College, Sattur (Affiliated to Madurai Kamaraj University) Madurai, Tamil Nadu, India Email:revathy7284@gmail.com



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ABSTRACT

The drug Salbutamol aids in the lungs medium and broad airway enlargement. Treatment for acute episodes of bronchospasm caused by bronchial asthma, chronic bronchitis, and other long-term bronchopulmonary disorders including chronic obstructive pulmonary disease (COPD) usually involves its use. In this work, the degree-based and degree splitting topological indices of salbutamol are computed using both a Python software and manual calculation. Furthermore, QSAR analysis of the topological indices are discussed using SPSS

Keywords: Salbutamol, Topological indices, Degree Splitting graphs, Python Programming, SPSS

INTRODUCTION

A topological index of graph G is a numerical value that describes its topology. It reflects the theoretical characteristics of a chemical molecule when applied to its molecular structure. In this study, chemical structures of drugs used to treat asthma were subjected to well-known degree-based topological indices. Chemical structure is viewed as a graph, where the constituents are the vertices and the boundaries between them are the edges. Salbutamol is a member of the group of drugs known as bronchodilators, more specifically, 2-adrenergic agonists. This medicine is used to treat and prevent bronchospasm caused by respiratory illnesses like chronic bronchitis, asthma, and other breathing problems.





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Preliminaries

In this section, certain well-known definitions and findings concerning various topological indices of graphs are described for quick reference while one reads the material presented in this paper.

Definition 2.1:The degree of a vertex v in G or simply d(v) is the number of edges of G incident with vertex v.

Definition 2.2: ABC (atom bond connectivity) index of a graph G defined in [2] as, ABC(G) (atombond connectivity) index of a graph G in as,

$$ABC(G) = \sum_{pq \in E(G)} \sqrt{\frac{d(u) + d(v) - 2}{d(u)d(v)}}$$

Definition 2.3: ABS (atom bond sum connectivity) index of a graph G defined

in [1] as ABS(G) =
$$\sum_{pq \in E(G)} \sqrt{\frac{d(u)+d(v)-2}{d(u)+d(v)}}$$

Definition 2.4: ABS (atom bond sum connectivity) index of a graph G defined in [1] $\operatorname{asAZI}(G) = \sum_{pq \in E} {}_{(G)} \left(\sqrt{\frac{d(u)d(v)}{d(u)+d(v)-2}} \right)^3$

Definition 2.5:SAI (sum augmented index) is defined in [4] as,

$$SAI(G) = \sum_{pq \in E(G)} \left(\sqrt{\frac{d(u) + d(v)}{d(u) + d(v) - 2}} \right)^{3}$$

Definition 2.6: GA (geometric-arithmetic index)of a graph G is defined in [8]as

$$\mathrm{GA}(\mathrm{G}) \ = \ \sum\nolimits_{pq \in E \ (\mathrm{G})} \frac{2\sqrt{d(u)d(v)}}{d(u) + d(v)}$$

Definition 2.7:AG (Arithmetic-geometric index) of a graph G is defined in [8]as

$$AG(G) = \sum_{pq \in E(G)} \frac{d(u) + d(v)}{2\sqrt{d(u)d(v)}}$$

Definition 2.8:GO1(first Gourava index) and GO2 (second Gourava index) of a graph G are defined in [5] as,

$$GO1(G) = \sum_{u \in E(G)} \left[\left(d(u) + d(v) \right) + d(u)d(v) \right]$$

$$GO2(G) = \sum\nolimits_{pq \in E \ (G)} \left[\left(d(u) + d(v) \right) d(u) d(v) \right]$$

Definition 2.9:HGO1(first hyper Gourava index) and HGO2 (second hyper Gouravaindex)of a graph G are defined in [6] as,

$$HGO1(G) = \sum_{pq \in E(G)} (d(u) + d(v) + d(u)d(v))^{2}$$

 $HGO2(G) = \sum_{v \in E(G)} (d(u) + d(v)d(u)d(v))^{2}.$

Definition 2.10 : Let G=(V,E) be a graph with $V=S_1US_2....UT$ where each S_i is a set of vertices having at least two vertices , having the same degree and $T=V\setminus US_i.The$ Degree splitting of Graph denoted by DS(G) is obtained from G by adding new vertices w_1 , $w_2...w_i$ and joining w_i to each vertex of S_i





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Topological Indices of Salbutamol

The several topological indices of salbutamol are covered in this section. Salbutamol has 17 vertices and 17 edges in its chemical graph. The graph below displays salbutamol's chemical structure.

Theorem 1: ABC index of Salbutamolis ABC(G) = 12.6348

Proof: From definition 2.2

$$\begin{split} & \text{ABC(G)} \ = \ \sum_{pq \in E \ (G)} \sqrt{\frac{d(u) + d(v) - 2}{d(u)d(v)}} \\ & = \ 1 \sqrt{\frac{1 + 2 - 2}{(1)(2)}} \ + \ 2 \sqrt{\frac{1 + 3 - 2}{(1)(3)}} \ + \ 3 \sqrt{\frac{1 + 4 - 2}{(1)(4)}} \ + \ 2 \sqrt{\frac{2 + 2 - 2}{(2)(2)}} \ + \ 6 \sqrt{\frac{2 + 3 - 2}{(2)(3)}} \ + \ 1 \sqrt{\frac{2 + 4 - 2}{(2)(4)}} \ + \ 2 \sqrt{\frac{3 + 3 - 2}{(3)(3)}} \end{split}$$

= 12.6348

By using Python Programming, the ABC $\,$ of Salbutamol is also calculated.

```
#ABC(G)
import math
foriinrange(1,8):
a=int(input("enter a value:"))
b=int(input("enter b value:"))
if a==1and b==2:
  c=(a+b-2)/(a*b)
  print(c)
  c1_ans=math.sqrt(c)
  mul1=c1_ans*1
  print(mul1)
if a==1and b==3:
  c=(a+b-2)/(a*b)
  print(c)
  c2_ans=math.sqrt(c)
  mul2=c2_ans*2
  print(mul2)
 if a==1and b==4:
  c=(a+b-2)/(a*b)
  print(c)
  c3_ans=math.sqrt(c)
  mul3=c3 ans*3
   print(mul3)
if a==2and b==2:
  c=(a+b-2)/(a*b)
   print(c)
  c4_ans=math.sqrt(c)
  mul4=c4_ans*2
  print(mul4)
```





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```
if a==2and b==3:
  c=(a+b-2)/(a*b)
  print(c)
  c5_ans=math.sqrt(c)
  mul5=c5_ans*6
  print(mul5)
if a==2and b==4:
  c=(a+b-2)/(a*b)
  print(c)
  c6_ans=math.sqrt(c)
  mul6=c6_ans*1
  print(mul6)
if a==3and b==3:
  c=(a+b-2)/(a*b)
  print(c)
  c7_ans=math.sqrt(c)
  mul7=c7_ans*2
  print(mul7)
final=mul1+mul2+mul3+mul4+mul5+mul6+mul7
print(final)
```

Output of the above program

```
enter a value:1
enter b value:2
0.7071067811865476
enter a value:1
enter b value:3
0.666666666666666
1.632993161855452
enter a value:1
enter b value:4
0.75
2.598076211353316
enter a value:2
enter b value:2
0.5
1.4142135623730951
enter a value:2
enter b value:3
0.5
4.242640687119286
```





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enter a value:2 enter b value:4 0.5 0.7071067811865476

12.635470518407578

Similarly, the other topological indices from Definition 2.3 to 2.9 are calculated by both analytically and using Python programming which are listed in the table below:

Topological Index	Topological Index of Salbutamol
ABC	12.6354
ABS	12.8266
AZI	116.6424
SAI	110.7916
GA	15.782
AG	18.3044
GO1	250
GO2	458
HGO1	7248
HGO2	15572

Topological Indices of Degree Splitting of Salbutamol

The different topological indices of Salbutamol's degree splitting are assessed in this section. The degree splitting graph of salbutamol consists of 21 vertices and 34 edges.

The topological indices from Definition 2.2 to 2.9 are calculated by both analytically and using Python programming which are listed in the table below:

Topological Index	Degree Splitting Topological Index of Salbutamol
ABC	22.6959
ABS	28.9427
AZI	433.2746
SAI	91.6532
GA	30.8491
AG	36.2013
GO1	1011
GO2	3360
HGO1	48111
HGO2	388584

Comparison Between Salbutamol & Degree Splitting Graph of Salbutamol

This section presents the correlation coefficient, as determined by a regression model, between Salbutamol's topological indices and its degree splitting graph. Using the SPSS software and the data in Tables 1 and 2, the linear and quadratic models are produced.





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Best-fit values

Slope	22.42 ± 2.696
Y-intercept	-9224 ± 14649
X-intercept	411.4
1/Slope	0.04460

95% Confidence Intervals

Slope	16.20 to 28.64
Y-intercept	-43005 to 24558
X-intercept	-1281 to 1776

Goodness of Fit

R square	0.8963
Sy.x	41640

Is slope significantly non-zero?

F	69.18
DFn,DFd	1,8
P Value	< 0.0001
Deviation from horizontal?	Significant

Data

2	
Number of XY pairs	10
Equation	Y = 22.42*X - 9224

CONCLUSION

Salbutamol's degree based topological indices and degree splitting topological indices are examined using degree-based indices. A graphical comparison of the computed findings for the chemical compounds stated above may be found in Figure 1. The creation of novel drugs to treat chronic bronchitis, bronchial asthma, and other long-term bronchopulmonary conditions may benefit from these discoveries.

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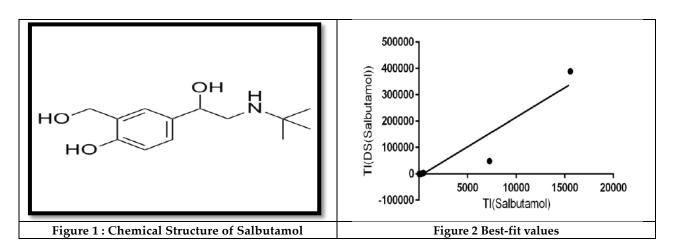
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Table 1. Edge partition of Salbutamol

$(d(u),d(v))/(u,v)\inE(G)$	Number of edges
(1,2)	1
(1,3)	2
(1,4)	3
(2,2)	2
(2,3)	6
(2,4)	1
(3,3)	2

Table 2. Edge Partition of degree Splitting of Salmabutamol

$(d(u),d(v))/(u,v)\in (G)$	Number of edges
(1,5)	1
(2,3)	2
(2,4)	2
(2,5)	3
(2,6)	6
(3,3)	2
(3,4)	5
(3,5)	1
(3,6)	6
(4,4)	6







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RESEARCH ARTICLE

The Impact of COVID 19 Pandemic on the Professional Learning and Evaluation of Undergraduate Pharmacy Students

Srujankumar Reddy Pothula¹, Sravani Jollireddy², Dipak Dnyandeo Bharambe³ and Mohanraj Rathinavelu4*

¹Pharmacist, Department of Pharmacy, Bharat Heavy Electricals Limited Organisation (Electronics Division) Karnataka, India.

²Assistant Professor, Department of Pharmacy Practice, Oil Technological and Pharmaceutical Research Institute, (Affiliated to JNT University) Anantapur, Andhra Pradesh, India.

³Assistant Manager, Department of Clinical Pharmacy, P D Hinduja Hospital and Medical Research Center, Mumbai, Maharashtra, India.

⁴Associate Professor, Department of Pharmacy Practice, Raghavendra Institute of Pharmaceutical Education and Research (RIPER) (Affiliated to JNT University) Anantapur, Andhra Pradesh, India.

Received: 30 Dec 2023 Revised: 19 Jan 2024 Accepted: 27 Mar 2024

*Address for Correspondence Mohanraj Rathinavelu

Associate Professor,

Department of Pharmacy Practice,

Raghavendra Institute of Pharmaceutical Education and Research (RIPER)

(Affiliated to JNT University) Anantapur,

Andhra Pradesh, India.

Email: visitmoley@gmail.com



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ABSTRACT

In the higher education teaching-learning process, evaluation is extremely important, it aids in the formation of judgmental values, educational status, and student accomplishment. The global pandemic of the Coronavirus has produced major change, bringing numerous difficulties to the higher education community. After four months in the midst of the global crisis, we've realized that the COVID-19 is here to stay, and we need to find ways to move forward. This crisis might be viewed as an opportunity to rebuild our long-standing educational systems and establish better, more up-to-date academic methods that are appropriate for today's learners. In this context, the current study investigated the perceptions of pharmacy students to the teaching and learning process in pharmacy education at undergraduate level, towards blended pedagogy by administering a standard validated questionnaire; and the educational outcomes with examination as indicators conducted as open-book examination and closed-book examination assessed the outcome through the results of end semester examination at undergraduate level in a self-financing, unaided, private pharmacy institute of south India. All the observations in our study were prolific in terms of teaching learning process through blended learning among students. In





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addition, we observed that the end semester examination outcome is satisfactory in in terms of pass percentage, which had direct correlation with individual student outcome in terms of gender almost in most of cases. In conclusion, our study envisaged effective student engagement, with more facilitator-student interactions and adaptability; through blended learning which enabled, enhanced and transformed students to active learners.

Keywords: Assessment, Blended Pedagogy Coronavirus disease 2019, Open Book Examination, Pharmacy Students.

INTRODUCTION

The health, social and economic crises of the COVID-19 pandemic have thrown into sharp relief the disruptive forces acting on the global higher education (HE) sector [1,2]. With more than 3 million COVID 19 infection cases as of August 2020, India experienced an accelerated uptake of digital solutions, tools, and services, speeding up the transition towards digital education [3]. During the coronavirus crisis, the education sector has gone through a rapid transformation, Universities adopted e-learning was a cure-all to COVID-19 [4]. Considering the national scenario; India's apex regulatory body of higher education, University Grants Commission (UGC), mandated for all the universities to complete the 25% syllabus through online teaching mode and 75% face-to-face interaction (UGC, 2020) [5]. The COVID 19 pandemic have stalled the pharmaceutical education which highly pressed the priority to review, re imagine, and revisit the Indian pharmaceutical education which underwent a didactic change including the implementation of novel strategies of online education and complete curriculum revision by the Pharmacy Council of India (PCI). Across the globe blended learning is applied in pharmaceutical education, but it gained momentum in Indian pharmacy education during the global threat COVID 19 pandemic[6]. Therefore, for successful implementation of educational change implications of change need to be addressed with blended pedagogy enabling a greater access to course content, learning preparation in peers and interactions [7].

Additionally, the outcome of teaching-learning process is dependent and proportional to the third pillars in higher education which is called evaluation [8]. In Indian higher education at university level adapted open book examination (OBE) for students' evaluation but was not considered as gold standards. However, self-financing private institutes was not able to cope up with this OBE due to technical shortcoming, finance and student's perception. The sudden closure of institutions/universities has not only adversely impacted the learning of the students worldwide; unfortunately, the school closure situation coincided with the key assessment period of the students, and their exams were either postponed or have been cancelled that year. As a result, parents and teachers fail to assess a student's learning capabilities which can have long-term negative consequences for him or her. The studies conducted in the past have shown that while comparing students' scores for OBE and closed book examination (CBE), there was no statistical difference, even though students who took CBE had a slightly higher score. Hence, the current study was performed to investigate the perceptions of pharmacy students towards blended pedagogy in pharmaceutical education and outcome of teaching learning process during COVID 19 and post COVID 19 through examinations at undergraduate level in a self-financing, unaided, private pharmacy institute of south India.

MATERIALS AND METHODS

Research design and study

The current research applies a quantitative design where descriptive statistics are used for the student characteristics (age and gender variables), and features data to determine if they are significant in blended learning effectiveness and outcome evaluation.





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This study is based on an experiment (online survey Google Form administered) in which learners participated during their on-line session (COVID 19) of B. Pharmacy program course teaching during academic year 2019-2020.

Research Instrument

The online self-regulated learning questionnaire [9] and the intrinsic motivation inventory [10] were adopted and applied to measure the constructs on self- regulation in the student characteristics and motivation in the learning outcome constructs. Other self-built instruments were used to study remaining variables (attitudes, computer competence, workload management, social and family support, satisfaction, knowledge construction, technology quality, and interactions).

Instrument reliability

Cronbach's alpha [11] (measure of internal consistency, that is, how closely related a set of items are as a group, test for reliability), one of the most important and pervasive statistics in research involving test construction and use [12]; was applied used to test reliability. All the scales and sub-scales had acceptable internal consistency reliabilities are within range and found to be significant within acceptable values, ranging from 0.70 to 0.95 [13-15].

Study participant

The experimental survey was performed online, by administering the validated questionnaire prepared through google form to students pursuing undergraduate program (B. Pharmacy) during the academic year 2019-2020 at an unaided self-financing private pharmacy institute (Raghavendra Institute of Pharmaceutical Education and Research RIPER Autonomous) in rural Andhra Pradesh and the responses were documented and analyzed in the Microsoft spread sheet.

Study procedure

The current quantitative online experimental survey of six months duration was performed in an unaided self-financing private autonomous pharmacy institute of rural Andhra Pradesh state. The study included students of both genders, pursuing undergraduate (B. Pharmacy) pharmacy program at the institute respectively, who showed willingness towards participation. Ethical approval was obtained from the institute before the start-up of study by submitting a detailed proposal with potential outcome.

Preparation of a validated questionnaire

A questionnaire was prepared to investigate the perception of students at undergraduate level towards blended pedagogy in pharmacy education by using the literatures [9,10] available in the core area.

Pilot study

A preliminary pilot pretesting of the questionnaire (comprising of 39 inventories) was done on 20 each randomly selected students from both undergraduate programs, with the purpose of identifying the practical and communication difficulties while surveying. Furthermore, the pilot testing allowed us to modify the ambiguous and unsuitable questions.

Assessment of reliability and internal consistency

The reliability and internal consistency of the questionnaire based on Cronbach's alpha coefficient value [12] were between 0.7 and 0.953 [13-15], and was found reliable for 28 inventories.

Final draft and standard

Furthermore, the questionnaire after its final drafting was reviewed by expert panel in subjects and curriculum for the face validity, content validity, and the relevance and comprehensiveness Thus, the validated questionnaire comprised of 28 inventories under 7 categories (online resources, interactions, quality of technology, self-management, students approach towards blended learning, motivation and satisfaction), along with the demography particulars of the students. Furthermore, the same validated questionnaire was administered to





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students pursuing undergraduate program (B. Pharmacy) at our institute through Google Form due to the COVID 19 pandemic and lockdown in the academic year of 2019-2020, within the duration of February to June, 2020; who showed willingness towards the study. The responses were presented in 5-point Likert scale based on satisfaction, and the same was collected and analyzed using Microsoft spread sheet.

Program outcome evaluation (COVID 19 and post COVID 19)

It was observed that during COVID 19 pandemic, the institute (RIPER Autonomous) have convened two sessional examinations (I and II) in open book examination (OBE) mode and end semester examination in closed book examination (CBE) for undergraduate program (B. Pharmacy) students of academic year (AY) 2019-2020. The results of which, obtained from examination cell are collected and collated, and further evaluated for defining the outcomes as pass percentage in terms of course and program.

RESULTS AND DISCUSSION

The current study performed in an unaided private self-financing autonomous pharmacy institute of rural Andhra Pradesh to investigate the perception of pharmacy students towards blended pedagogy in pharmacy education, through an online experimental survey among students pursuing undergraduate pharmacy program (B. Pharmacy) during the academic year AY 2019-2020 for a period of six months. The study was performed by administering a validated self-administered questionnaire online, among a total of 501 undergraduate students (B. Pharmacy) at the institute through social networking applications.

Characteristics of study participants

Out of 448 administration, 430 respondents filled the form, which observed 2.79% of incomplete information, so the total completed responses obtained was 418. In our undergraduate student population of 418, 218 (52.15%) were female, and 200 (47.85%) were male. During the academic year AY 2019-2020 out of 418 complete responses, students of I year II semester 112 (with 41.07% male and 58.93% female) were more in comparison to students of other year and semesters, where students of II and III years were same in total (24.88% each) relatively equal gender distribution, observations of which is presented in Table 1.

Response towards online administration of validated self-administered questionnaire

Blended learning courses may be implemented in several ways[16,17]. Three categories described by Graham[16] cover a range of possibilities: enabling blends, enhancing blends, and transforming blends. Our study, observed an enhanced student perception towards online resources in terms of use ableness (68.6%), content (74.8%), and practical worth (62.5%); were facilitator purposely blended the teaching-learning process in to the course of the program, which involved a combination of face-to-face interactions with small group activities, through both synchronous and asynchronous computer-mediated interactions; and practice-oriented learning activities. Through which traditional classroom limitations were overcome, which transformed the passive learning environment to active learning environment. As discussed earlier the pedagogy facilitated by the teacher gained a didactic momentum, and were nothing out- of -box discussion was made, with face-to-face interactions and small group activities. Thus, high rate of teaching and learning experience was experienced by students through facilitation of student-faculty (78.88%) and student content interactions (70.4%), all the time during the course class, interactions was important to achieve enhanced blending. Blended learning approaches are being applied in pharmacy education programs across the learning continuum[18-20], but this have been adopted in India too due to the COVID 19 pandemic, which made the teaching learning process so interesting for students, with considerable limitations. The same was observed in our study, despite three-fourth of students experience quality materials and contents (75.4%) and enhanced face-to-face support (68.3%), the regular accessibility to technology was observed in one-half of the students, due to the rural and low-income backdrop of students. The learner satisfaction factor (self-management) is of high priority in blended learning, which provides them an opportunity to make a self-assessment, and identify their potentials and enables capacity building in terms of setting of strategies (72.7%), critical thinking (66.9%), time





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management (75.8%) and maintaining an optimistic learning environment (68.81%). The students of our study (more than two-third on average) were able to map this clearly, showcasing a higher rate of satisfaction with blended pedagogy, build through an effective self-assessment (69.5%). Literature[21] suggests that there is a significant relationship between academic success and learner autonomy; academic success and motivation; and learner autonomy and motivation in terms blended learning (BL) environments. Blended learning refers to asynchronous distance mode learning coupled with face-to-face facilitation of pedagogy, whose success is highly influenced by student's autonomy (72.4%), interactions (78.8%), family and societal support (71.4%), and the standards of academic or course instructions (74.6%), our study observed all these indicators at highest level, with a considerably more workload (56.46%) which may not influence the success of blended learning, in turn it builds up stress among students. Hence, the facilitator has to take a heightened responsibility in guiding them in management of stress whatsoever it matters. The study observed a two-third satisfaction on an average in terms of all the indicators which influence the blended pedagogy (teacher 71.8%, course content 74.8%, technology transfer 58.7%, interactions 78.8%, and constructive knowledge 73.7%). In our study, all the observations were prolific in terms of teaching learning process through blended learning (reaction, learning, and behavior) among students, but the actual effectiveness of blended learning prevails on the extent of achieving the program outcomes, which has to be visualized through their performance in examinations, results of which are reported in Figure 1.

Matrix of course and outcomes

Evaluation has been supplied by C.E. Beeby (1977) [22] who described evaluation as "the systematic collection and interpretation of evidence leading as a part of process do a judgement of value with a view to action." Evaluation is an important component of the teaching-learning process. It helps teachers and learners to improve teaching and learning. Evaluation is a continuous process not a periodic exercise. In our study, we observed that the end semester examination outcome is satisfactory in in terms of pass percentage secured in respective courses, which had direct correlation with individual student outcome in terms of gender (were female secured appreciable pass percentage) almost in most of cases. On the other hand, pass percentage in some courses observed was lesser in male students likely in courses pharmaceutical organic chemistry (46%), biochemistry (54%), medicinal chemistry (55.22%), biopharmaceutics and pharmacokinetics (50.82%), and biotechnology (55.74%) respectively; which was highly influenced by internal examination results convened through open book examination, results of which are presented in Figure 2. This pandemic has made evaluation process handicapped for higher education and put us in wheel chair, though university grant commission (UGC) have suggested OBE has a mode of examination, the experience in Indian higher education is not enough prepared due to slow adoption to digitization and less equipped technocracy and utilization both in teachers and students. On a microscopic observation, student of our institute appears for online examination during the regular course of study in B. Pharmacy program; and the same was followed in midterm examination during pandemic where the outcome is good in almost all individual cases but less in few, findings of which is similar to other investigated studies. [23] The consideration of few may be due to their cognition as complicated production of questions and marking schemes, subjective marking and restriction of materials. In our experience and from our observations; when properly used, open-book examination is a beneficial assessment tool that proves to be popular among students because it eliminates the cueing effect, encouraging students to gain a deeper understanding of their learning material while also requiring the use of key skills such as critical thinking. We also strongly recommend effective evaluation process supports evident and productive decision-making process; which promotes teaching-learning process for which blended pedagogy is a gateway outweighing all complexities of pharmacy and pharmaceutical sciences teaching and learning (T and L) fulfilling classroom objectives.

CONCLUSION

In conclusion, our study envisaged effective student engagement, with more facilitator-student interactions and adaptability; through blended learning which enabled, enhanced and transformed students to active learners; and recognized that there are drawbacks to administering an online open book exam which prioritized further research in terms of efficacy and practicalities of online OBEs in Indian higher education.





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Table 1. Demography of study participants

Program	Year and Semester	Gender distribution		Total	
		Male	Female	n	%
B. Pharmacy	I & II	46 (41.07%)	66 (58.93%)	112	26.79
	II & II	47 (45.19%)	57 (54.81%)	104	24.88
	III & II	54 (51.92%)	50 (48.08%)	104	24.88
	IV & II	53 (54.08%)	45 (45.92%)	98	23.44
	Total	200 (47.85%)	218 (52.155)	418	100

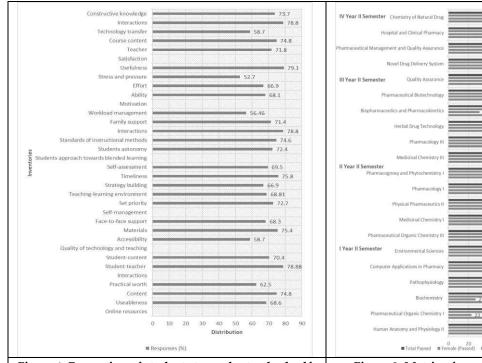


Figure 1. Perception of students towards standard selfadministered questionnaire on blended learning

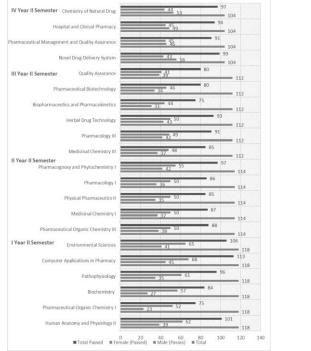


Figure 2. Matrix of program and course outcome





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RESEARCH ARTICLE

Assessing the Suitability of Groundwater for Irrigation in Pandavapura Taluk, Mandya District, Southern Karnataka, India

Madhu R1* and P Madesh2

¹Research Scholar, Department of Studies in Earth Science, University of Mysore, Karnataka, India. ²Professor, Department of Studies in Earth Science, University of Mysore, Karnataka, India.

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*Address for Correspondence Madhu R

Research Scholar, Department of Studies in Earth Science, University of Mysore, Karnataka, India. Email: madhu.r.geo@gmail.com



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ABSTRACT

The groundwater suitability for irrigation in Pandavapura Taluk's comprehensive investigation employs a diverse set of hydrochemical indices to assess the quality of groundwater, including the Permeability Index, Salinity and Sodium Hazard Classification, Sodium Adsorption Ratio (SAR), Residual Sodium Carbonate (RSC), Sodium Ratio, Kelly's Ratio and Magnesium Adsorption Ratio (MAR). The research focuses on evaluating the groundwater's potential for sustainable agricultural use by examining key parameters related to water quality. The findings from this comprehensive analysis will provide valuable insights for farmers, policymakers, and water resource management authorities in Pandavapura Taluk, Mandya district.

Keywords: Irrigation, Groundwater, Pandavapura, water resource management, Mandya

INTRODUCTION

Water stands out as the most widely accessible and cost-effective raw material. Its diverse applications derive from its distinct physical and chemical characteristics. India's available water resources are insufficient to adequately irrigate the cultivatable land. As a result, it is essential to make concerted efforts to optimize the utilization of water resources for agricultural irrigation. Agriculture constitutes the primary livelihood for numerous communities in semi-arid regions, in Pandavapura taluk where agriculture is of paramount importance and its water supply relies on a combination of groundwater and a non-perennial river. To ensure successful crop growth and productivity, agriculture in these regions heavily depends on supplementary irrigation. However, the distribution of river water across the area is uneven. Therefore, groundwater serves as a primary irrigation water source due to its greater





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reliability compared to surface water sources. The monsoons play a vital role in replenishing the water resources during June to October, as well as excessive extraction during the pre-monsoon period from February to May. Over-exploitation of groundwater, a common practice in many semi-arid areas, can lead to the depletion of aquifers and result in land subsidence, thereby diminishing both the availability and quality of groundwater. The chemical composition of irrigation water can directly impact plant growth through either toxicity or nutrient deficiencies, and indirectly by altering the accessibility of nutrients to plants. Irrigation water typically contains various dissolved salts. In the absence of proper management, considering the potential adverse effects on plants and soils, prolonged utilization of such water sources becomes unfeasible. This study aims to evaluate the suitability of groundwater for irrigation in the specified area.

Study Area

The study area, Pandavapura, lies between approximately 76°36' to 76°38' East longitude and 12°25' to 12°43' North latitude (Fig. 1), covering 529.9 square kilometers. It is depicted on topographical maps 57D/10, 57D/11, and 57D/14. Summers extend from February to May, while winters span mid-November to mid-February, the primary rainy season is from June to September with temperatures ranging from 16°C to 35°C. The average annual rainfall of about 650 millimeters. The area boasts a diverse composition of rock types, including migmatites, granodiorite tonalite gneiss, granite, metaultramafite, schist, metabasalt, and tuff from Archean to Proterozoic eras.

METHODS AND METHODOLOGY

The Pre-monsoon groundwater samples were collected from hand pumps and borewells across a systematically divided 10x10 square kilometer grid for comprehensive coverage. A total of 45 samples were carefully collected in clean polythene bottles, labeled, and borewells were pumped for 5-10 minutes to minimize iron pipe influence. GPS coordinates were recorded for accurate mapping in ArcGIS 10.8. Key parameters like TDS, pH, and EC were promptly measured in the field. In the lab, calcium, bicarbonate, carbonate, and chloride were assessed through titration, while calcium was determined by EDTA titration. Sodium and potassium were quantified using a flame photometer, and sulfate, fluoride, iron, and nitrate levels were determined through spectrophotometry. Data underwent processing in WATCHIT software, and thematic maps were generated using ArcGIS.

RESULT AND DISCUSSION

Groundwater Chemistry

The hydrochemical data for the pre-monsoon seasons are shown in Table.1. The pH range of 5.53 to 8.11 is attributed to natural geological features, atmospheric CO₂, microbial activity, human influence, and other environmental factors (Hitchon et al., 1999). Electrical conductivity (EC) reflects dissolved ions and salinity, affected by aquifer leaching, saline sources, or a combination (Hem 1991; Hounslow 1995). Pre-monsoon groundwater EC levels in this taluk range from 208 to 1600 μ S/cm, while Total Dissolved Solids (TDS) vary from 105 to 689 mg/L, mainly comprising inorganic salts. Major ions influencing hydrochemical traits include Ca²⁺, Mg²⁺, Na⁺, Cl, SO₄², HCO₃, K, and NO₃, with concentrations falling within specific ranges (Li et al., 2016). Concentrations of Ca²⁺, Mg²⁺, and Na⁺ fall within the ranges of 24.8 to 237.6 mg/L, 6.8 to 92.34 mg/L, and 13.1 to 666 mg/L, respectively. Chloride (Cl) concentrations vary from 25.8 to 241.48 mg/L, while sulfate (SO₄) ranges from 1.8 to 432.3 mg/L. Bicarbonate (HCO₃) concentrations span from 43.7 to 735 mg/L, potassium (K⁺) levels range from 0.8 to 154 mg/L, and nitrate (NO₃) concentrations vary between 0.01 and 115 mg/L (Table 1).

Irrigation Water Quality Parameter

The assessment of water suitability for irrigation is influenced by specific water quality standards established by various organizations and authorities (Ayers R 1994). Irrigation of flat land primarily depends on groundwater as its main source. The quality of irrigation water is influenced by a variety of factors, including soil composition, soil





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texture, local geology, and agricultural practices (Abou El-Defan et al.,2016). Some factors that determine the quality of water used for irrigation are Kelly's Ratio, Magnesium Adsorption Ratio, Permeability Index, Salinity, Sodium Adsorption Ratio, Soluble Sodium Percentage and Sodium Ratio (Kshitindra Kr Singh et al., 2020).

Salinity Hazard Classification

Salinity Hazard Classification assesses and categorizes soil salt levels, crucial for agriculture. High salinity from saline irrigation reduces plant water absorption, harming crop growth and yield. If unaddressed, it leads to significant crop yield loss (Abou El-Defan et al., 2016). In 1954, Richards categorized irrigation water salinity into four groups, each with a letter grade (Table 2&3). This system sets thresholds for salinity hazard levels and guides decisions on soil management and mitigation strategies for land managers and agricultural practitioners. In the research area only 1 sample proved suitable for irrigation, whereas nearly 41 samples raised doubtful and three samples were deemed unfit for irrigation.

Sodium Ratio (SR)

The Sodium Ratio (SR), which is determined by comparing the concentration of dissolved sodium cations to the combined levels of calcium and magnesium ions in the irrigation water (Abou El-Defan et al., 2016). The sodium level is determined using the equation provided below.

$$SR = (Na^+)/(Ca^{2+} + Mg^{2+})$$

In good water, this ratio should not exceed one (Table 4). All ion concentrations are expressed in meq/L. In the specified research area, only 10 out of the 45 samples were classified as inappropriate, while the remaining 35 samples were deemed suitable for agricultural purposes Fig.(2,3).

Soluble Sodium Percentage (SSP)

Sodium's vital for classifying irrigation water due to its effect on soil, reducing permeability. Na+ percentage gauges water suitability for irrigation (Wilcox, 1955). Groundwater quality is assessed based on Na% as follows: >20% (excellent), 20 - 40% (good), 40 - 60% (permissible), 60 - 80% (doubtful), and >80% (unsafe) (Table 5). The Na+% are determined through a formula, in which all ionic concentrations are expressed in meq/l.

$$Na\% = (Na^+ + K) / Ca^{2+} + Mg^{2+} + Na^+ + K) \times 100$$

Among 45 samples, 8 are excellent, 25 good, 5 permissible, and 1 unsafe Fig. (4).

Permeability Index (PI)

Soil permeability is influenced by sodium, calcium, magnesium, and bicarbonate levels, which also affect long-term irrigation water quality (Doneen 1964). Permeability is classified into three classes: I, II, and III (Table 6). The equation provided below is utilized to calculate the permeability index.

$$PI = \frac{Na + \sqrt{HCO_3}X100}{Ca + Mg + Na}$$

The concentrations of ions are uniformly expressed in meq/L. In the specified study area, 43 out of 45 samples were deemed suitable for agriculture, while the other two were unsuitable. Fig. (5).

Kelly's Ratio (KR)

Kelly's ratio (KR) assesses irrigation water suitability (Kelley 1963). A KR over one indicates unsuitability for irrigation (Aravinthasamy et al., 2020).

$$KR = \frac{Na}{(Ca + Mg)}$$

The Kelly's Ratio in the study region indicates that 35 samples were unsuitable, while the remaining 10 samples were within the acceptable range. (Table 7) & Fig. (6).

Sodium Absorption Ratio (SAR)

The SAR is crucial in evaluating water quality. Elevated salt levels in irrigation water can raise soil sodium levels, impacting permeability and causing water infiltration issues, leading to soil dispersion and cultivation challenges





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(Keesari et al., 2016). SAR is calculated by comparing the sodium concentration to the square root of the calcium and magnesium concentrations in the water sample (Uma Mohan and Krishnakumar, 2021). The subsequent equation is employed for its calculation.

$$SAR = \frac{Na}{\sqrt{\frac{Ca + Mg}{2}}}$$

These concentrations are typically measured in units of meq/L. The USSL diagram was created to assess water quality Fig. (7, 8), Out of 45 samples, 37 were excellent, 5 good, 2 doubtful, and 1 unsuitable for irrigation. (Table 8, 9)

Residual Sodium Carbonate (RSC)

Residual Sodium Carbonate (RSC) measures potential sodium-related soil issues from bicarbonate (HCO₃) and carbonate (CO₃²-) ions in irrigation water. It's computed by subtracting total calcium (Ca) and magnesium (Mg) ion concentrations from total bicarbonate and carbonate ions. A positive RSC signals excess sodium, leading to soil dispersion and reduced permeability during irrigation.

$$RSC = (CO_3^- + HCO_3^-) - (Ca^{2+} + Mg^{2+})$$

Negative RSC occurs when calcium and magnesium ions exceed bicarbonate and carbonate levels, indicating less sodium buildup in soil. This makes water suitable for irrigation and agriculture (Table 10). In the specified research area, 44 out of the 45 samples were classified as being in the good category, indicating their suitability for irrigation. Conversely, the remaining one sample fell into the poor category, signifying its unsuitability for irrigation. Fig. (09).

Magnesium Adsorption Ratio (MAR)

Calcium ions and magnesium ions are typically found in a natural equilibrium in the soil. However, this equilibrium can be disrupted on occasion when there is an elevated concentration of magnesium ions. An excess of magnesium ions can potentially hinder plant growth by increasing the alkalinity of the water. (Fallatah & Khattab, 2023). The calculation of the MAR was determined using a specific formula.

$$MAR = \frac{Mg X 100}{Ca + Mg}$$

The MAR in the study region indicates that 34 samples were suitable, while the remaining 11 samples were within the inappropriate for irrigation. (Table 11) & Fig. (10).

CONCLUSION

The suitability of groundwater for irrigation in semi-arid region pandavapura based on multiple parameters has yielded important insights. Sodium hazard classification results indicate that the majority of the samples fall into the doubtful category, with only a small percentage being classified as good. This classification suggests the need for careful monitoring and management to prevent potential harm to soil and crops. Furthermore, the sodium ratio and Kelly ratio results suggest that a significant portion of the groundwater samples is unsuitable for irrigation, emphasizing the importance of water quality management practices to mitigate the adverse effects on soil structure and crop growth. The Permeability Index highlights that the vast majority of samples (43 out of 45) are either highly suitable or suitable, demonstrating the water's ability to infiltrate and drain effectively in the soil. Additionally, the Sodium absorption ratio predominantly falls into the "excellent" category, which is an encouraging finding for irrigation practices. Residual sodium carbonate is generally good, with only a minor presence of "poor" samples, indicating acceptable conditions for irrigation. Finally, the Magnesium Absorption Ratio (MAR) results show that the majority of the samples are suitable for irrigation, although a portion falls into the "unsuitable" category. In summary, the groundwater in the study area exhibits a mixed suitability for irrigation, with certain parameters suggesting potential challenges. Effective management and monitoring practices, as well as the development of sitespecific irrigation strategies, will be essential to ensure the sustainable use of groundwater for agricultural purposes in the region. This study serves as a valuable resource for local farmers and policymakers to make informed decisions regarding irrigation practices and water quality management.





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Table 1: Statistical assessment of physicochemical parameters in groundwater.

Parameter	Unit	Min	Max	Mean	Std. Deviation
рН	ı	5.53	8.11	7.33	0.57
TDS	Mg/L	104	689	378.34	113.77
EC	μS/cm	208	1600	793.24	263.16
TH	Mg/L	122	852	388.51	141.06
SO ₄	Mg/L	1.8	432.3	53.02	63.84
F	Mg/L	0.1	2.74	1.0384	0.44975
Fe	Mg/L	0.02	0.29	0.083	0.080
Mg	Mg/L	6.8	92.34	36.68	19.05
Cl	Mg/L	25.8	241.48	74.94	44.80
NO ₃	Mg/L	0.01	115	18.88	26.63
HNO ₃	Mg/L	43.7	735	278.48	119.10
K	Mg/L	0.8	154	12.873	28.83
Ca	Mg/L	24.8	237.6	92.036	40.411
Na	Mg/L	13.1	666	126.05	146.75

Table 2: Classes of irrigation water salinity (Richards, 1954).

Tubic	Table 2. Classes of Hilgation water saminty (Menards, 1934).					
Class	Description	Salinity	T.D.S.	Remarks		
C1	Low Salinity	<25	<200	This irrigation method suits many crops and soils. Regular leaching is vital to remove excess salts, especially in poor or porous soils, which may require increased irrigation.		
C2	Medium Salinity	0.25- 0.75	200- 500	This irrigation technique suits areas with stable leaching, where soil salt content doesn't rise significantly. It's ideal for plants tolerant to moderate salinity, suitable for widespread cultivation without special salinity controls.		
C3	High Salinity	0.75-2.5	500- 1500	In areas with poor drainage, this irrigation method should not be used. Even if the drainage is adequate, special measures are needed to manage salinity, and only salt-tolerant plants can be grown.		
C4	Very High Salinity	>2.5	>1500	In most cases, this irrigation method should not be used. However, it may be used in special circumstances, such as when the soil is permeable, there is good drainage, and crops that can tolerate high salt levels are grown. In these cases, the irrigation water must be applied in excess to leach out salts.		

Table:3 Salinity Hazard (after Davis and De Weiest, 1966; Wilcox, 1955)

Salinity Hazard Classification	Remarks on quality	Number. of Samples	Percentage
C1	Excellent	0	0
C2	Good	1	2.2
C3	Doubt	41	91.1
C4	unsuitable	3	6.7

Table 4: Sodium ratio (SR) classification (after Abou- El-Defanet al., 2016)

:	Sl No.	Values meq/l	Remarks on quality	Numbers of Samples	percentage
	1	>1	Unsuitable for agriculture	10	22.2
	2	<1	Suitable for agriculture	35	77.8





Madhu and Madesh

Table 5: Soluble Sodium Percentage (SSP) classification (after Wilcox, 1954).

Category	S.S.P. Range (%)	No. of samples	Percentage
Excellent	<20	08	17.8
Good	20-40	25	55.6
Permissible	40-60	5	11.1
Doubtful	60-80	6	13.3
Unsafe	>80	1	2.2

Table 6: Permeability Index (Doneen, 1964)

Sl No.	Class	Remarks on quality	No of samples	Percentage
1	Class I (>75%)	Water is classified as having a high probability of providing irrigation benefits.	06	13.3
2	Class II (25- 75%)	Water is classified as having a high probability of providing irrigation benefits	37	82.2
3	Class III (<25%)	Water is unsuitable with a 25% overall permeability	02	4.4

Table 7: Kelly's Ratio (Kelly, 1940)

Kelly's Ratio	Remarks	No of samples	Percentage
>1	Suitable	10	22.2
<1	Unsuitable	35	77.8

Table 8: Irrigation water classes and description as SAR values (after Abou-El-Defanet al., 2016)

Class	Description	S.A.R. Value	Remarks
	-	(meq/l)	
S1	Low Sodium water	<10	This irrigation method is safe to use on most soils, as it does not cause sodium to build up in the soil. However, sodium-sensitive crops may accumulate harmful levels of sodium if this irrigation method is used.
S2	Medium Sodium water	10-18	Fine-textured soils with a high CEC are at risk of becoming saline, especially if they do not have gypsum and are not leached regularly. This irrigation method can be used on coarse-textured or organic soils with good permeability, but it is not recommended for fine-textured soils.
S3	High Sodium water	18-26	This irrigation water is generally not recommended, as it could lead to high levels of exchangeable sodium in the soil, which can be harmful to plants. Most soils will require special management practices to mitigate these risks, such as good drainage, high leaching, and the addition of organic matter. However, this water may be safe to use in gypsiferous soils, which contain the mineral gypsum. Gypsum can help to replace exchangeable





			sodium with calcium, which is less harmful to
			plants.
S4	Very High Sodium water	>26	This water is too saline for most crops, but it may be possible to use it for irrigation of salt-tolerant crops in soils with high calcium levels or where gypsum is added to the water

Table 9: Sodium Hazard classification (Richards, 1954)

Sodium Hazard Class	S.A.R. range meq/l	Remark	No. of Samples	Percentage
S1	<10	Excellent	37	82.2
S2	10-18	Good	5	11.1
S3	18-26	Doubtful	2	4.4
S4	>26	Unsuitable	1	2.2

Table 10: Residual Sodium Carbonate (after Abou-El-Defanet al., 2016)

Category	R.S.C. range (meq/l)	Water Quality	No. of samples	Percentage
Good	< 1.	Water probably safe for irrigation.	44	97.8
Medium	1.25-2.50	Water is marginally suitable for irrigation and can be used with certain conducts.	0	0
Poor	>2.50	Water is unsuitable for irrigation purposes.	1	2.2

Table 11: MAR Classification of the groundwater samples in the study area (after Gupta and Gupta, 1987)

MAR	Remarks	No of samples	Percentage
< 50	Suitable	34	75.6
>50	Unsuitable	11	24.4

Table 12: Parameters indicating the quality of irrigation water in the research area.

Sample No.	Location	Lattitude	Longitude	SR	SSP	RSC	USSL salinity
1	Chikkayarahally	12.4396	76.5866	0.44	32	-2.98	C3
2	Anthanahalli	12.4690	76.5436	0.41	31.3	-8.57	C3
3	Bindahalli	12.4852	76.5490	0.16	14.9	-12.01	C4
4	Maligere	12.5021	76.5338	0.07	7.3	-3.26	C3
5	Dinka	12.5198	76.5457	0.54	35.4	-1.53	C3
6	Aralakuppe	12.4582	76.6163	0.10	9.8	-4.29	C3
7	Seethapura	12.4457	76.6131	0.13	11.4	-2.09	C3
8	Chaluvarasanakoppalu	12.4362	76.6382	0.45	32	-1.04	C3
9	Kyatanahalli	12.4638	76.6527	0.27	22	-0.12	C2
10	Kennalu	12.4725	76.6750	1.41	65.7	0.15	C3
11	Pandavapura	12.5042	76.6734	1.84	65.4	-0.36	C3
12	Hulkere	12.5026	76.6206	0.47	32.2	-3.14	C3
13	Chinakuruli	12.5383	76.6036	1.48	59.9	-6.53	C3
14	Kurubaramanchanahalli	12.5746	76.5782	0.32	25.1	-6.2	C3
15	Morasanahalli	12.5560	76.5966	2.26	69.6	-3.03	C3
16	Hiremaralli	12.5200	76.6933	4.59	83.9	5.76	C3
17	Kenchnalli	12.5344	76.6355	0.13	11.7	0.85	C3





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18	Kerethanoor	12.5573	76.6443	1.97	67.6	-0.9	C3
19	Kodala	12.5928	76.6423	0.57	37	-2.98	C3
20	Marmalli	12.6156	76.6133	0.61	38.1	-4.01	C3
21	Kadaba	12.5873	76.6121	0.61	38.9	-1.01	C3
22	Gowdagere	12.6308	76.6641	0.65	39.8	-5.34	C3
23	Chikade	12.4977	76.6928	0.12	13	-0.6	C3
24	Talasasana	12.4879	76.7281	0.26	21	-4	C3
25	Dodda byadarahalli	12.5029	76.7464	0.51	34.4	0.83	C3
26	Agathahalli	12.5232	76.7800	1.33	57.2	-2.29	C3
27	Kanaganamaradi	12.5297	76.7294	0.47	34.2	-0.87	C3
28	Kurahatti	12.5568	76.7266	0.30	23.2	-3.59	C3
29	Kalenahalli	12.5690	76.7462	0.36	29.8	-4.18	C3
30	Madarahalli	12.5865	76.7278	0.32	25.5	-3.28	C3
31	Balenahalli	12.6315	76.7242	0.42	30.1	-1.63	C3
32	G.Hosahalli	12.6471	76.7264	0.55	36.2	-1.82	C3
33	Shamboonahalli	12.6377	76.7068	0.59	38.1	-0.42	C3
34	K.hosur	12.5389	76.6700	0.32	24.4	-4.09	C3
35	Madeshwarapura	12.5715	76.6796	1.47	60.1	-3.43	C3
36	Iakanahalli	12.6496	76.6809	0.46	31.7	-9.14	C3
37	Talekere	12.7045	76.6484	0.13	13.1	-12.69	C4
38	Melkote	12.6625	76.6505	0.47	32.6	-1.91	C3
39	Belatheguppe	12.5499	76.6902	3.33	78.2	-3.3	C4
40	Anunahalli	12.6071	76.6897	0.72	42.5	-2.69	C3
41	Basavanagudikopallu	12.5476	76.5668	0.73	42.6	-0.34	C3
42	Garudapura	12.6870	76.6744	0.08	7.6	-3.86	C3
43	shamboonahalli	12.4965	76.6464	1.05	51.3	-3.48	C3
44	Baby grama	12.4963	76.5899	0.41	29.4	-1.05	C3
45	Alphalli	12.4794	76.5935	0.24	20.9	-5.81	C3

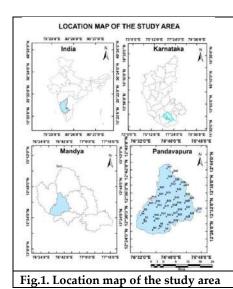
Table 13: Parameters indicating the quality of irrigation water in the research area.

Location	Lattitude	Longitude	SAR	KR	Na%	MAR	PI	Na hazard
Chikkayarahally	12.4396	76.5866	2.47	0.44	31.97	38.00	50.16	S1
Anthanahalli	12.4690	76.5436	2.98	0.41	31.33	56.93	40.57	S1
Bindahalli	12.4852	76.5490	1.21	0.16	14.91	48.59	23.39	S1
Maligere	12.5021	76.5338	0.48	0.07	7.34	45.99	30.87	S1
Dinka	12.5198	76.5457	2.56	0.54	35.44	35.52	58.32	S1
Aralakuppe	12.4582	76.6163	0.59	0.1	9.83	8.92	31.20	S1
Seethapura	12.4457	76.6131	0.70	0.13	11.44	46.90	38.61	S1
Chaluvarasanakoppalu	12.4362	76.6382	2.08	0.45	31.99	36.71	58.07	S1
Kyatanahalli	12.4638	76.6527	0.83	0.27	22.04	49.18	70.20	S1
Kennalu	12.4725	76.6750	6.71	1.41	65.67	40.50	76.22	S1
Pandavapura	12.5042	76.6734	7.19	1.85	65.36	38.95	81.99	S1
Hulkere	12.5026	76.6206	2.50	0.47	32.20	47.50	51.09	S1
Chinakuruli	12.5383	76.6036	9.49	1.48	59.90	54.45	67.28	S2
Kurubaramanchanahalli	12.5746	76.5782	1.78	0.32	25.09	53.90	36.57	S1
Morasanahalli	12.5560	76.5966	14.63	2.27	69.57	32.21	77.36	S3
Hiremaralli	12.5200	76.6933	23.06	4.59	83.91	38.03	91.95	S4





Kenchnalli	12.5344	76.6355	0.65	0.13	11.74	46.32	48.62	S1
Kerethanoor	12.5573	76.6443	9.98	1.97	67.61	42.92	78.60	S2
Kodala	12.5928	76.6423	3.25	0.57	37.00	24.29	54.14	S1
Marmalli	12.6156	76.6133	3.87	0.61	38.05	41.22	52.92	S1
Kadaba	12.5873	76.6121	2.69	0.61	38.94	37.27	62.98	S1
Gowdagere	12.6308	76.6641	4.45	0.65	39.84	33.51	52.55	S1
Chikade	12.4977	76.6928	0.59	0.12	13.04	33.61	44.99	S1
Talasasana	12.4879	76.7281	1.56	0.26	21.02	30.21	40.45	S1
Dodda byadarahalli	12.5029	76.7464	2.13	0.51	34.37	39.73	67.85	S1
Agathahalli	12.5232	76.7800	8.07	1.33	57.24	50.49	69.33	S2
Kanaganamaradi	12.5297	76.7294	2.00	0.47	34.19	46.09	60.39	S1
Kurahatti	12.5568	76.7266	1.64	0.3	23.19	25.81	43.07	S1
Kalenahalli	12.5690	76.7462	2.19	0.36	29.78	39.34	44.45	S1
Madarahalli	12.5865	76.7278	1.82	0.32	25.51	19.43	44.83	S1
Balenahalli	12.6315	76.7242	2.30	0.42	30.07	38.64	52.59	S1
G.Hosahalli	12.6471	76.7264	2.35	0.55	36.16	52.21	58.92	S1
Shamboonahalli	12.6377	76.7068	2.65	0.59	38.13	25.86	64.20	S1
K.hosur	12.5389	76.6700	2.00	0.32	24.38	25.63	42.49	S1
Madeshwarapura	12.5715	76.6796	7.32	1.47	60.12	52.17	70.27	S2
Jakanahalli	12.6496	76.6809	3.08	0.46	31.69	59.63	40.37	S1
Talekere	12.7045	76.6484	0.97	0.13	13.14	11.62	17.23	S1
Melkote	12.6625	76.6505	1.92	0.47	32.58	52.38	56.38	S1
Belatheguppe	12.5499	76.6902	17.63	3.33	78.23	43.00	83.24	S3
Anunahalli	12.6071	76.6897	3.42	0.71	42.48	46.14	59.35	S1
Basavanagudikopallu	12.5476	76.5668	2.65	0.73	42.63	16.92	72.15	S1
Garudapura	12.6870	76.6744	0.43	0.08	7.62	53.15	30.79	S1
shamboonahalli	12.4965	76.6464	6.81	1.05	51.35	29.04	63.57	S2
Baby grama	12.4963	76.5899	2.04	0.41	29.39	51.04	54.76	S1
Alphalli	12.4794	76.5935	1.38	0.24	20.87	50.06	34.52	S1



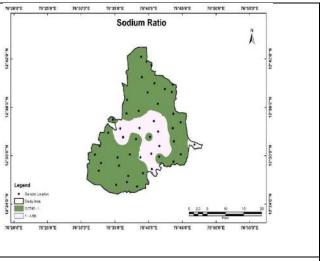
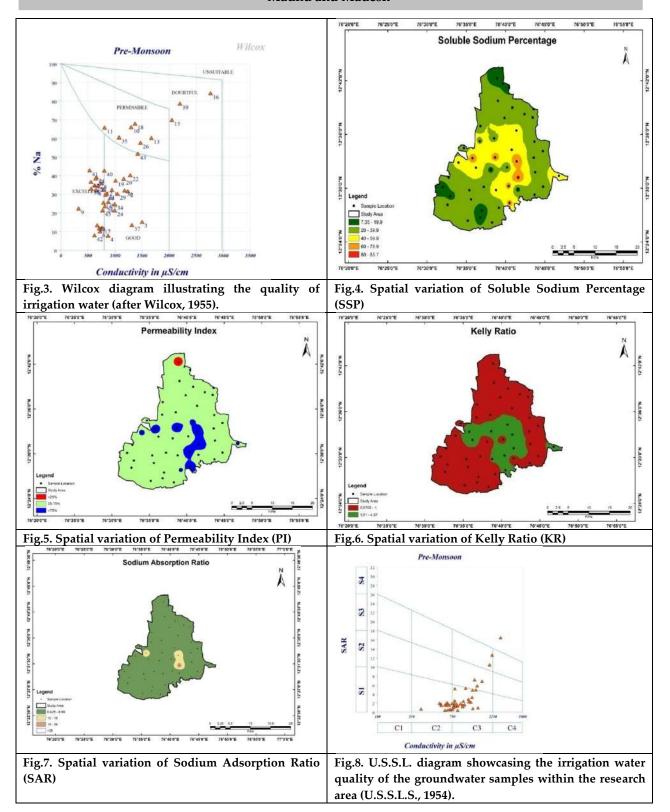


Fig.2. Spatial variation of Sodium Ratio

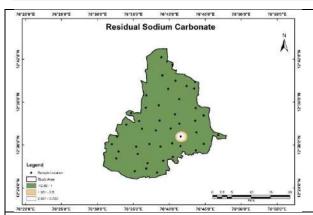












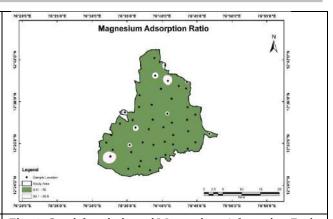


Fig.9. Spatial variation of Residual Sodium Carbonate (RSC)

Fig.10. Spatial variation of Magnesium Adsorption Ratio (MAR)





RESEARCH ARTICLE

Functional Recovery Following Early Strengthening After Middle Third **Clavicle Fracture-A Single Case Study**

Veluri Rakesh 1*, Arunachalam Ramachandran2 and Anandhan Duraisamy3

¹Ph.D Scholar, Madhav College of Physiotherapy, Faculty of Medical Sciences, Madhav University, Pindwara, Sirohi, Rajasthan, India.

²Professor, Madhav College of Physiotherapy, Faculty of Medical Sciences, Madhav University, Pindwara, Sirohi, Rajasthan, India.

³Associate professor, Madhav College of Physiotherapy, Faculty of Medical Sciences, Madhav University, Pindwara, Sirohi, Rajasthan, India.

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*Address for Correspondence Veluri Rakesh

Ph.D Scholar, Madhav College of Physiotherapy, Faculty of Medical Sciences, Madhav University, Pindwara, Sirohi, Rajasthan, India. Email: dr.rakeshphysio91@gmail.com



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ABSTRACT

Clavicle fractures are common injuries that can result in prolonged pain, or non-union, inadequate functional movements, if treated improperly. Often the rehabilitation gets delayed due to various reasons which when attested a better functional recovery may be attained. The objective of the report is to narrate novel implementation of early strengthening program within the period of six weeks for middle third clavicle fracture. The patient in this report, a 21 years old male, who had a middle third clavicle fractures, treated with early strengthening program followed by a surgical management showed convincing recovery in pain and function.

Keywords: Clavicle fracture, early strengthening, Rehabilitation.

INTRODUCTION

The clavicle is the most often broken bone in the human body, accounting for up to 5% to 10% of all fractures. [1,2] These injuries are especially common in younger patients and are frequently related with direct clavicle trauma, such as in contact sports and motor vehicle accidents. Males are more affected than females, and the prevalence decreases with age, yet traumatic falls in older patients generate a bimodal peak in the age distribution. [2]Fractures are the





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most prevalent, which accounts for about 80% of all clavicle fractures.[3, 4]The gold standard is plate and screw fixation, often known as open reduction and internal fixation [ORIF]. There is a surgical alternative. Plate fixation has the advantage of being less technically demanding than intramedullary fixation. [5] Plate mending research has provided promising outcomes. They claimed a unionization rate of 97% and a no unionization rate of 94%. [6] The rate of postoperative satisfaction in 232 midshaft clavicle fractures. They discovered that ORIF significantly decreased non-union and complication rates when compared to non-operative treatment for displaced midshaft clavicle fractures. [7] This study's main objective was to determine whether an early start-up strengthening program within six weeks would speed up functional recovery in a comparable manner to post-surgical treatment.

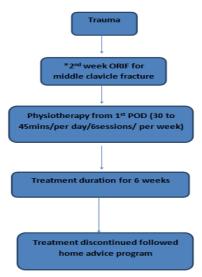
Patient Information

On November 27, 2022, a male, age 21, was involved in a bike accident. The patient was admitted to a nearby General Hospital due to loss of consciousness. Further medical examination was performed because the patient's vital signs were unstable and he was vomiting continuously. Due to such unfavourable medical condition, the underlying fracture was not diagnosed initially and the patient was managed conservatively for two weeks. Then he was admitted to an orthospeciality hospital, where he was diagnosed with a left medial one-third clavicle comminuted fracture through X-ray [Figure 1A]. The patient was successfully treated with open reduction and internal fixation [Figure 1B] and was discharged on the fifth post-operative day.

Physiotherapy Assessment

Following surgery, the patient was sling-immobilized advice his arm adducted for approximately one month. The Numerical Pain Rating Scale [NPRS] was initially used to assess the pain severity, which was 8/10. There was edema, left deltoid muscle atrophy, abrasions on the left forearm and hand, and a depressed shoulder with the arm adducted on examination. Palpatory examination revealed that the scar was adherent and had grade 1 tenderness. The active and passive range of motion and muscle power of the shoulder were slightly compromised during the examination using bubble inclinometer and push pull dynamometer, Functional outcome of an individual was measured using quick dash score[shown in table 1 & 2]. The patient reports difficulty with daily tasks such as lifting heavy objects, bathing, and overhead activities.

Time Line



^{*}Delayed due to unfavourable condition.





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Diagnostic Assessment

Show the figure 1 A] X-ray showing the mid third fracture. B]-X-ray Showing the open reduction with internal fixation[ORIF]

Therapeutic Intervention

StandardMedical management like medications, suturing, and wound care were offered by professionals. The main goal of physiotherapy sessions was to enhancement is to strengthen the shoulder muscles and daily tasks. The operative extremity was placed in a sling for two weeks immediately following surgery, so early active resisted range of motion of the elbow, wrist, and hand was encouraged. Intervention was offered for 30 to 45 minutes of one session/day, 6 sessions per week for 6 weeks.

Weeks 1-2:Daily 3 to 5 minutes of soft tissue mobilisation across the suture area; active elbow, wrist, and hand range of motion; and active shoulder range of motion with active assistance. The passive 90° flexion/abduction range of motion of the shoulder joint was encouraged.

Weeks 3 to 4:Soft tissue mobilisation across the suture region for 3 to 5 minutes. Internal and external shoulder flexion, abduction, and rotation exercises. Perform strengthening exercises for the elbow, wrist, and hand movements with dumbbells. Pendular and a wandexercise, Workouts with pulleys, ladders, and mariners wheel are encouraged.

Weeks 5 and 6:Scapular strengthening workout using dumbbells and theraband followed by scapula stabilisation exercises. Strength and functional training should be increased in preparation for a gradual return to activity.

DISCUSSION

Most previous studies have recommended sling immobilization and physical therapy following operative treatment of a clavicle fracture, using arm sling protection for two to six weeks after that followed by rehabilitation.[8,9,10]A Randomized controlled trail was done to compare open reduction and plate fixation versus non operative treatment for displaced mid shaft clavicular fracture. The results reveal that open reduction with internal fixation is expensive and associated with implant related complications, which are absent in non-operative treatment. They conclude that non operative treatment is more feasible choice for treating midshaft clavicular fracture. [7] A study was done by Karibasappa A G and Srinath S.R to find out the effectiveness of surgical versus conservative treatment for managing displaced midshaft clavicular fracture, they concluded that the group which received open reduction internal fixation with osteosynthesis showed good results as compared to the other group which received non-surgical treatment.[11]. Physiotherapy plays a vital part in healing quickly and returning to everyday activities by using fixation, immobilisation by braces in the first week, and gentle exercise in the sixth week following injury, followed by active exercise and muscular strength after 12 weeks.[12]

The main limitation of the current study is the lack of an objective tool to mark the recovery and the lack of long-term follow up. These aspects should be attested through robust research methods in future. Physical therapy is started as much as possible from the day one of the postoperative care in this trial, and it includes scar management and early strength training to improve the patient's ability to function. Scar management includes scar massage and soft tissue mobilisation. Early strengthening began after six weeks in a previous study, but in the current study, it began in the third week of the exercise regimen. As a result, early strength training will help to promote functional activities in the individual as soon as possible.

Patient Perspective

With a fractured collarbone, I've been in a lot of pain and under a lot of stress. My level of confidence has increased while receiving treatment, which helps me get through pain and return to my functional tasks.





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Informed Consent

The patient agreed to provide his written, fully informed consent and his permission to record his thoughts on the course of treatment.

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Table 1 -Pre and Post Outcomes for NPRS and Quick Dash disability

Outcome	1st day	3 rd week	6th week
Numerical pain rating scale [NPRS]	8/10	4/10	2/10
Quick dash Disability	80%	40%	20%

Table 2 -Pre And Post Range Of Motion [ROM AND MMT]

	LEFT SHOULDER										
MOVEMENT	ACTIVE ROM				PASSIVE ROM			MANUAL MUSCLE TESTING [MMT]			
	1 st	3rd	6 th	1 st	3 rd	6 th	1 st	3 rd	6 th		
	Day	Week	Week	Day	Week	Week	Day	Week	Week		
FLEXION	60	110	170	80	120	180	2/5	4/5	5/5		
EXTENSION	10	15	20	15	20	25	2/5	3/5	5/5		
ABDUCTION	40	90	140	45	110	170	2/5	3/5	5/5		
INTERNAL	20	50	75	25	70	90	2/5	3/5	5/5		
ROTATION	20	30	75	23	70	70	2/3	3/3	3/3		
EXTERNAL	10	35	50	15	60	75	2/5	3/5	5/5		
ROTATION	10	33	30	13	00	7.5	2/3	3/3	3/3		





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Figure 1. A] X-ray showing the mid third fracture. B]-X-ray Showing the open reduction with internal fixation [ORIF]





RESEARCH ARTICLE

GC/MS Analysis of Methanolic Extract of Pancratium triflorum Roxb. bulbs

Rakkimuthu R1*, Neethu Krishna K2, Sathishkumar P3, Ananda Kumar AM3 and Sowmiya D3

¹Assistant Professor and Head, PG and Research Department of Botany, Nallamuthu Gounder Mahalingam College, Pollachi (Affiliated to Bharathiar University), Coimbatore, Tamil Nadu, India. ²PG Student, PG and Research Department of Botany, Nallamuthu Gounder Mahalingam College,

³Assistant Professor, PG and Research Department of Botany, Nallamuthu Gounder Mahalingam College, Pollachi (Affiliated to Bharathiar University), Coimbatore, Tamil Nadu, India.

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*Address for Correspondence Rakkimuthu R

Assistant Professor and Head, PG and Research Department of Botany,

Nallamuthu Gounder Mahalingam College, Pollachi

(Affiliated to Bharathiar University), Coimbatore, Tamil Nadu, India.

Pollachi (Affiliated to Bharathiar University), Coimbatore, Tamil Nadu, India.

Email: biorakki@gmail.com



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ABSTRACT

This study presents the chemical composition of the methanolic extract obtained from Pancratium triflorum bulbs through gas chromatography and mass spectrometry analysis. Six distinct compounds were identified, namely 2,3-Anhydro-D-Galactosan, Triacontanoic Acid, Methyl Ester, 1-Hexadecyne, Eicosonoic Acid, Cis-9,10 Epoxyoctadecan-1-OL, and 2-Isopropyl-5-Methylcyclohexyl 3-(1-(4-Chlorophenyl)-3-Oxobutyl)-C. This comprehensive profiling provides insights into the molecular constituents of Pancratium triflorum, contributing to our understanding of its chemical composition and potential pharmacological applications. Further investigations into the biological activities of these identified compounds may unveil novel therapeutic possibilities associated with Pancratium triflorum.

Keywords: Pancratium triflorum, Bulb, GC-MS, Bioactive compounds, medicinal plant

INTRODUCTION

Pancratium triflorum. Roxb, a perennial herbaceous plant native to the Indian subcontinent, has a long-standing history of traditional use in various indigenous medicinal systems. Its cultural importance is underscored by its botanical nomenclature in which 'Pancratium' signifies 'all strength' and 'triflorum' refers to its characteristic threeflowered inflorescence [1]. In traditional practices, Pancratium triflorum is celebrated for its therapeutic attributes,





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including its anti-inflammatory, diuretic, and analgesic properties [2]. However, despite its well-documented ethnobotanical uses, there exists a paucity of comprehensive scientific investigations exploring the phytochemical constituents and biological properties of this plant. Recent advancements in analytical techniques have provided the means to scrutinize the chemical composition of medicinal plants, thus shedding light on their pharmacological potential. Gas Chromatography-Mass Spectrometry (GC/MS) stands out as a powerful tool for the identification of bioactive compounds within plant extracts. This study employs GC/MS analysis to decipher the complex phytochemical profile of *Pancratium triflorum*. This research seeks to fill a significant knowledge gap and provide a solid foundation for further exploration of the medicinal and therapeutic potential of *Pancratium triflorum*.

The Amaryllidaceae family, to which *Pancratium triflorum* belongs, is well-known for the synthesis of alkaloids with diverse pharmacological properties. This is exemplified by the presence of lycorine, a prominent alkaloid known for its anti-inflammatory, cytotoxic, and antiviral activities, which is found in various Amaryllidaceae species [3]. Furthermore, the family Amaryllidaceae includes several species that have displayed notable antioxidant potential, which aligns with the increasing scientific interest in natural antioxidants as potential alternatives to synthetic antioxidants due to their lower toxicity and broader health benefits [4]. In light of the aforementioned, this research delves into the GC/MS analysis of *Pancratium triflorum* to uncover its phytochemical constituents. The outcomes of this study could pave the way for further investigations into the therapeutic applications of *Pancratium triflorum*. Roxb. and the development of antioxidant-based therapies.

MATERIALS AND METHODS

Plant Material and Extraction

Pancratium triflorum. bulbs were collected from karimkulam, palakkad district kerala. The plant material was identified and authenticated by botanical Survey of India, Coimbatore, and a voucher specimen (NGMPGB/20) was deposited in a herbarium. The bulbs were cleaned, shade-dried, and powdered for extraction

Preparation of Plant Extract

Powdered plant material (50 g) was w subjected to soxhlet extraction using methanol for 72 hours at room temperature. The extract was then filtered and concentrated under reduced pressure to obtain a crude methanolic extract of *Pancratium triflorum*.

GC-MS Analysis

The GC/MS analysis was performed using the Clarus 680 GC was used in the analysis employed a fused silica column, packed with Elite-5MS (5% biphenyl 95% dimethylpolysiloxane, 30 m \times 0.25 mm ID \times 250 μ m df) and the components were separated using Helium as carrier gas at a constant flow of 1 ml/min. The injector temperature was set at 260°C during the chromatographic run. The 1 μ L of extract sample injected into the instrument the oven temperature was as follows: 60 °C (2 min); followed by 300 °C at the rate of 10 °C min–1; and 300 °C, where it was held for 6 min. The mass detector conditions were: transfer line temperature 230 °C; ion source temperature 230 °C; and ionization mode electron impact at 70 eV, a scan time 0.2 sec and scan interval of 0.1 sec. The fragments from 40 to 600 Da. The spectrums of the components were compared with the database of spectrum of known components stored in the GC-MS NIST [5] (2008) library.

RESULTS AND DISCUSSION

The GC/MS analysis of the methanolic extract of *Pancratium triflorum*. revealed the presence of bioactive compounds. A total of 6 compounds were identified (Table 1). Notable compounds included carbohydrate derivatives, fatty acid derivatives, aliphatic hydrocarbons, eicosanoid derivatives, and structurally complex compounds. The Gas Chromatography-Mass Spectrometry (GC/MS) analysis of the methanolic extract of *Pancratium triflorum*. has unveiled a diverse array of bioactive compounds, contributing to the understanding of the plant's potential





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pharmacological properties. In this study, a total of six compounds were identified, major compounds were Cis-9,10 Epoxyoctadecan-1-OL (39.240%) and 2-isopropyl-5-methylcyclohexyl 3-(1-(4-chlorophenyl)-3-oxobutyl)-c (35.494%) and minor compounds were shown in figure 1, each possessing distinctive chemical structures and, possibly, associated biological activities. These compounds are integral components of the plant's phytochemical composition and merit further investigation for their potential therapeutic applications.

Eicosonoic Acid, Cis-9,10 Epoxyoctadecan-1-OL, is a derivative of eicosanoids, which are signaling molecules involved in inflammation and immune responses [6]. The presence of this compound suggests that Pancratium may possess components with immunomodulatory properties. The compound 2-isopropyl-5methylcyclohexyl 3-(1-(4-chlorophenyl)-3-oxobutyl)-c, a complex and structurally diverse compound, adds further intrigue to the phytochemical profile of Pancratium triflorum. Its structure implies the potential for multifaceted pharmacological activities, which may encompass anti-inflammatory, antioxidant, or even cytotoxic effects. One of the identified compound, 2,3-anhydro-d-galactosan, is a carbohydrate derivative. While carbohydrates are not typically associated with bioactive compounds, their presence in plant extracts could indicate potential interactions with other bioactive molecules, and they may play a role in the overall medicinal properties of the plant. Triacontanoic Acid, methyl ester, is a long-chain fatty acid derivative. Fatty acids and their derivatives are known for their various biological activities, including anti-inflammatory and antimicrobial properties [7]. This compound could contribute to the plant's anti-inflammatory potential. 1-Hexadecyne is an aliphatic hydrocarbon. Aliphatic hydrocarbons are commonly found in plant extracts and can exhibit diverse biological activities, including antioxidant properties [8]. This compound may contribute to the antioxidant potential of Pancratium triflorum. While these compounds have been identified, their specific pharmacological properties and the potential synergistic interactions among them remain to be explored. Additionally, the concentrations of these compounds in the plant extract and their bioavailability are essential factors that need further investigation.

CONCLUSION

The GC/MS analysis of *Pancratium triflorum*. revealed the presence of several bioactive compounds, including carbohydrate derivatives, fatty acid derivatives, aliphatic hydrocarbons, eicosanoid derivatives, and structurally complex compounds. This diverse range of compounds suggests the potential for a broad spectrum of pharmacological properties. Further research is required to elucidate the specific roles of these compounds and their potential for contributing to the medicinal and therapeutic properties of *Pancratium triflorum*. Roxb.

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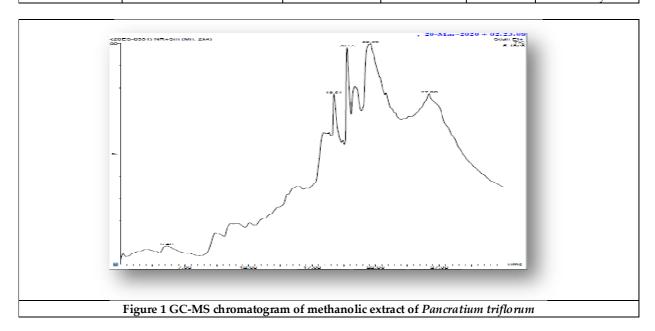




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T	able 1: Compounds identified	in the methanolic	extract of Pancra	atium trifloru	m
RT	Name of the compound	Molecular formula	Molecular weight	Peak area %	Compound nature
18.765	2,3-ANHYDRO-D- GALACTOSAN	C6H8O4	144	5.567	Aldehyde
19.495	TRIACONTANOIC ACID, METHYL ESTER	C31H62O2	466	6.154	Fatty acid methyl ester
20.566	1-HEXADECYNE	C16H30	222	5.774	Aliphatic hydrocarbon
21.171	EICOSANOIC ACID	C20H40O2	312	7.771	Unsaturated fatty acid
22.411	CIS-9,10- EPOXYOCTADECAN-1-OL	C18H36O2	284	39.240	Alcoholic compound
27.008	2-ISOPROPYL-5- METHYLCYCLOHEXYL 3- (1-(4-CHLOROPHENYL)-3- OXOBUTYL)-C	C30H33O6C1	524	35.494	Benzopyrone Antioxidant, antimicrobial and anti- inflammatory activity







RESEARCH ARTICLE

Novel Method of Controlling the Self Focusing and De-Focusing Length by using Non-Linear Kerr Material and Electro-Optic Materials

Rupali Maji

Assistant Professor, Department of Physics, New Alipore College (Affiliated of University of Calcutta), Kolkata, West Bengal, India.

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*Address for Correspondence Rupali Maji

Assistant Professor,

Department of Physics,

New Alipore College (Affiliated of University of Calcutta),

Kolkata, West Bengal, India. Email: rpl.maji@gmail.com

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ABSTRACT

Nonlinear optical materials are used for several physical applications. In optical switching, lens less optical focusing and defocusing these non-linear materials can show its strong applications. The focal length of a material (if the material is used for self-focusing) depends on the applied power. Here in this chapter, a method of controlling the focal length of a nonlinear material based on the joint use of electrooptic material and a nonlinear crystal is proposed. The focal length of the nonlinear material depends upon the voltage applied to the electro-optic material. By changing this voltage/or field in the electrooptic material, the focal length can be varied and this technique can be used as a focal length controller. A suitable electro-optic material and a nonlinear material can be used this purpose.

Keywords: Nonlinear optical materials, field, voltage, controller

INTRODUCTION

When the refractive index of a material is depends on the applied electric field linearly is called Pockel effect and when it depends on the square of the applied field is called Kerr effect. The Kerr effect also called quadratic electrooptic effect [1]. The Kerr effect has a distinct from the Pockel's effect as has the induced index change is directly proportional to the square of the electric field instead of varying linearly with it. This refractive index variation is responsible for the nonlinear optical effects like self-focusing, self-phase modulation and modulation instability and is the basis for Kerr-lens mode locking. There are several applications of Kerr effects in optical switching, arithmetic and algebraic operations etc. [2, 3, 4, 5, 6, 7, and 8]. In the Kerr electro-optic effect, or DC Kerr effect, a slowly varying external electric field is applied across the sample material. Under the influence of the external signal the sample birefringent, with different indices of refraction for light polarized parallel or perpendicular the applied field. The





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difference in index of refraction, Δn , $(\Delta n = n - n_0)$, where n and n_0 the refractive indices of the material with and without the applied of the external electric field respectively) is given by $\Delta n = \lambda KE^2$, where λ is the wavelength of the applied light, K is a material constant, and E is the strength of the electric field. This difference in index of refraction helps the material to act as a wave plate when the polarization of light is perpendicular of the applied electric field. If the material is kept between two 'crossed' linear polarizer's, no light come out when electric field is turned off, and almost all the light is transmitted for the application of the optimum value of the electric field. A higher value of the Kerr constant allow as a good transmission with a smaller applied electric field. In this particular work, the author shows by the use of Kerr material how one can control the focal length here the focal length of a non-linear material actually controlled by applied voltage, and the system is behaves like an optical lens.

Self-focusing and De-focusing of a Gaussian beam by the use of non-linear material

Due to a Kerr type of lensing, an intense optical pulse propagating in a non-linear medium experiences a self-focusing, where the beam diameter is decreased compared to of a weaker pulse. The physical mechanism is based on a Kerr nonlinearity with positive χ^2 . In this situation, the higher optical intensities of near to the beam axis, as compared to the off axis intensity, causes an increased refractive index in the inner part of the beam. This modified refractive index distribution acts like a focusing lens. The effect, occurring in the case of a negative χ^2 nonlinearity, self-defocusing, where a reduced refractive index is seen on the beam axis. A Kerr non-linear process which arises in a media exposed to intense electromagnetic radiation, and which produces a variation of the refractive index n as described by the formula n=n₀+n₂I, where n₀ and n₂ are the linear and non-linear components of the refractive index respectively, and I is the intensity of the light passing through it. The intensity distribution is taken spatially Gaussian, and the sign of the non-linear correction n₂ be either positive or negative, for self-focusing and defocusing [1, 9]. If the non-linear correction term n₂ is positive then in peripheral region the plane wave front takes a concave shape in the direction of the beam and is focused at the optical axis of the medium (Fig-1a).On the other hand if the n₂ is negative than central part of the beam goes faster than that of the peripheral region. Consequently, the plane wave front takes the shape of a convex shape direction of propagation and. Thus it defocused into the axis (Fig-1b).

Electro-optic material as an Amplitude modulator

Electro-optic modulator is an optical device in which an electrical signals exploiting the electro-optic effect and is used to modulate a proper beam of light. The modulation may be used to change the phase, frequency, amplitude, or polarization of the modulated beam. Modulation bandwidth at the gigahertz range is possible with the use of a laser based coherent controlled modulators. [10,11,12,13,14]. Certain materials change their optical properties when they are exposed to an electric field. This is caused by the forces that distort the positions and orientations of the molecules the material. The electro-optic effect gives the change in the refractive index from low frequency electric field to high one up to new range[15,16,17,18,19,20]. Some electric-optic materials are massively used as amplitude modulator such as Potassium di-deuterium phosphate (KD*P), Beta-barium borate (BBO), also Lithium niobate (LiNb03),Lithium Tantalite(LiTaO3) and Ammonium dihydrogen phosphate (NH4H2PO4,ADP) etc. In addition to these there are also some organic types of special polymer modulators. A schematic diagram of LiNb03 based electro-optic modulator is shown in the fig 2.

Guassian beam

Gaussian beam has its transverse electric field and intensity distribution which is well approximated by Gaussian functions. Many lasers emits beams that has a Gaussian profile, for that reason the laser is said to be operating on the fundamental transverse mode, or " TEM_{00} mode" in the laser's optical resonator. When this beam is refracted by a diffraction-limited lens, a Gaussian beam is transformed into another Gaussian beam [13,14].

The beam profile of a Guassian beam is shown in fig 3. An integrated scheme of controlling the self-focusing length of a bulk non-linear medium by the use of electro-optic material

The refractive indices of the electro-optic modulator is [3.9]





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$$n_{x'} = n_0 - \frac{1}{2} n_0^3 r E_Z$$

(1)

$$n_{y'} = n_0 + \frac{1}{2} n_0^3 r E_Z \tag{2}$$

Where 'r' is the material material constant. Ez, is the applied field along z direction.

First a linearly polarized wave polarized along the x-direction and (χ' is one of the bi-axial direction of the Electro-optic material) traveling along the z-direction through electro-optic material is considered (fig 4), We have applied an external electric field Ez along the Z-direction (C-axis), then the output wave at Z= ℓ (where ℓ is the length of the electro-optic material along z) would be given by

$$\varepsilon_{X'}(\ell) = \varepsilon_{X'}(0) \exp\{i[\omega t - n_0(\omega/c)\ell + (\omega/2c)n_0^3 r E_z \ell\}$$
(3)

Here $\mathcal{E}_{x'}(z)$ and $\mathcal{E}_{y'}(z)$ are the X' and Y' components of the electric field of the used light.

In a similar manner, a beam polarized along the Y-direction (where Y' is the other bi-axial direction of Electro-optic modulator) the output wave at $Z=\ell$ will be given by

$$\varepsilon_{v'}(\ell) = \varepsilon_{v'}(0) \exp \{ [\omega t - n_0(\omega/c)\ell + (\omega/2c)n_0^3 r E_z \ell \}$$
(4)

Now consider an incident wave polarized along the y direction is taken then it can be decomposed into two linearly polarized waves along two orthogonal direction as X' and Y' as these two components will have equal amplitudes and will be in phase Z=0, i.e; at the input of the crystal. Thus the two components which were in phase at Z=0 now develop a phase difference which is a function of the applied electric field (Ez). Thus the retardation at Z= ℓ between the two components will be

$$\gamma = (\omega/c) n \omega^3 r_{c3} Ez \ell = \omega n \omega^3 r_{c3} V/c$$
 (5)

Where V=Ez ℓ is the voltage applied across the crystal. One can define the 'half wave'

Voltage V_{π} as the voltage required to develop a phase shift of π between the two orthogonal polarization components So, $\gamma = \pi = (\omega/c) n \sigma^3 r \omega V_{\pi}$ (6)

Or $V_{\pi} = \lambda_0/2n_0^3 r_{63}$

Substituting the values of ε_X and ε_Y given by the equations (1) and (2), the expression of the total field (ε) becomes

$$\varepsilon = \frac{1}{2} \operatorname{Aexp}\{i[\omega t - (n_0/c)\omega \ell + (\omega/2c)n_0^3 r_{63} \operatorname{Ez} \ell]\}[1 - \exp(-i\gamma)]$$
(7)

Where
$$\gamma = (\omega/c) n_0^3 r_0^3 E_z \ell = \pi (V/V_{\pi})$$
 (8)

Thus the intensity of the output beam is given by

$$I_{0} = \frac{1}{2} \operatorname{Re}[\varepsilon \varepsilon^{*}] = \frac{1}{2} \operatorname{A}^{2} \sin^{2} \frac{1}{2} \gamma$$

$$= \frac{1}{2} \operatorname{A}^{2} \sin^{2}[(\omega/2c)n_{0}^{3}r_{63}V]$$
(9)

Where V=EzL is the applied voltage. The intensity of the input beam (Ii) is given by

$$I = \frac{1}{2} A^2 \tag{10}$$

ThusI₀/I_I=sin²(
$$\frac{1}{2}\pi V/V_{\pi}$$
) (11)

I₀/I_i is the transmission coefficient of the electro-optic modulator.

If Vis very less than V_π





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The I₀/I_≈
$$\left[\frac{\pi^2}{4} \frac{V^2(t)}{V_{-}^2}\right]$$
 (12)

Again the nonlinear refractive index of the Kerr type crystal is

$$n=n_0+n_2I_0$$
 . (13)

Putting the values of I₀ from equation (3.12) in equations (3.13)

n=n₀+n₂I_i
$$\left[\frac{\pi^2}{4} \frac{V^2(t)}{V_{\pi}^2}\right]$$

n-n₀=n₂I_i[
$$\frac{\pi^2}{4}\frac{V^2(t)}{{V_{\pi}}^2}$$
]

$$\Delta n = n_2 \text{Li} \left[\frac{\pi^2}{4} \frac{V^2(t)}{V_{-}^2} \right]$$
 (14)

We know the focal length (Lsf) of the Gaussian beam in a non-linear medium can be expressed as[1]
$$Ls = a \sqrt{\frac{n_0}{2\Delta n}}$$
 (15)

where a is the radius of the beam[1] (fig-5)

Now, putting the value of Δn from equation (14) we can get,

$$L_{sf} = a \sqrt{\frac{n_0}{2n_2I_i \frac{\pi^2}{4} \frac{V^2(t)}{V_{\pi}^2}}}$$
 (16)

It is known that input intensity is Ii , where

I=E02 (E0 is the amplitude of the electric field strength of the light at the time of introduction in the modulator of the

Thus the expression of the focal length can be written as,

$$L_{s} = \frac{aV_{\pi}}{\pi V(t)} \sqrt{\frac{2n_0}{n_2 E_0^2}} \tag{17}$$

$$L_{sf} = \frac{2V_{\pi}}{\pi V(t)} \frac{a}{E_0} \sqrt{\frac{n_0}{2n_2}}$$
 (18)

Now E(r) can be written in a radial function (where r is the radial position in the circular beam)

as
$$E(r) = E_0 \sqrt{1 - \frac{r^2}{a^2}}$$
 (19)

For the mean value of the energy flux density, we obtain the expression

 $<S>=v\epsilon E^2/2=[\epsilon CE_0^2/(2n_0)](1-r^2/a^2)$

$$=(\varepsilon_0 n_0 C E_0^2/2)(1-r^2/a^2) \tag{20}$$

The energy flux of the beam is given by

$$P = \int_{\sigma} \langle S. \rangle d\sigma = \pi \varepsilon_0 n_0 c E_0^2 \int_0^a (1 - r^2 / a^2) r dr$$
 (21)

Where σ is the cross-sectional area of the beam. After integration the total energy flux (P) can be written as $P=\pi\epsilon_0 n_0 c E_0^2 a^2/4$ (22)







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$$E_0 = \sqrt{\frac{P^2}{\pi \varepsilon_0 n_0 ca}} \tag{23}$$

Thus the self focal length

$$L_{s} = \frac{V_{\pi}}{\pi V(t)} \frac{a^2 \sqrt{\pi \varepsilon_0 n_0} c}{\sqrt{P}} \sqrt{\frac{n_0}{2n_2}}$$
(24)

RESULT

The result obtained in equation (24) can be used to obtain the focal length in Carbon bi sulphide, or in any other non-linear medium. For Carbon bisulphide, $n_0=1.62$, $n_2=0.22*10^{-19}m^2/w^2$, thus for a power P=10MW and beam radius a=1 cm,((the LiNbo3 is used as Electro-optic modulator before the non-linear material) ℓ_{sf} become 6.92m (considering $V_{\pi}=64V$ and V=64v also in equation (3.24))). Again if the applied voltage V=640V in LiNbO3the ℓ_{sf} becomes 0.692 m.

CONCLUSION

From the above analytical treatment it is seen that if an electro-optic Pockel cell is used before a Kerr-cell which extends the self-focusing then one can easily control the focal length of the self-focusing system by applying the desired amount of voltage at the electro-optic material. Similarly the defocusing length can also be controlled by the same mechanism. The whole scheme may extend a tremendous application in optical communication through optical fiber. These mechanisms can help the coupling of desired amount of light intensity in an optical fiber from a source in case of data communication. To use it in the application domain one can use a suitable electro-optic material and a suitable simple Kerr non-linear medium.

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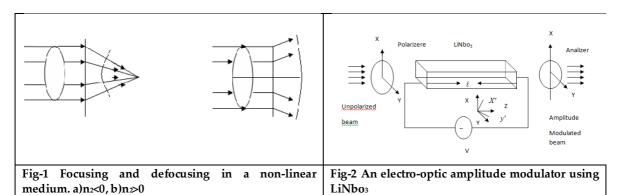
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Rupali Maji Intensity I Position x Fig-3 Profile of a Gaussian beam Fig-4 Scheme of controlling the focal length of Gaussian beam by use of E-o modulator and a Kerr type of non-linear material Ist Fig-5 Calculation of self-focusing length.





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RESEARCH ARTICLE

Effect of Cold therapy for Reduction of Perineal Pain among Postnatal Mothers in Selected Hospitals, Assam

Bhagya Priya Phukan^{1*}, Nongmeikapam Monika² and Zulanbeni C Kithan³

¹Student(M.Sc Nursing), Department of Obstetrics and Gynaecological Nursing, Faculty of Nursing, Assam down town University, Assam, India.

²Professor, Department of Obstetrics and Gynaecological Nursing, Faculty of Nursing, Assam down town University, Assam, India.

³Lecturer, Faculty of Nursing, Assam down town University, Assam, India.

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*Address for Correspondence

Bhagya Priya Phukan

Student(M.Sc Nursing),

Department of Obstetrics and Gynaecological Nursing,

Faculty of Nursing,

Assam down town University,

Assam, India.

Email: nangbhagyapriyaphukan@gmail.com



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ABSTRACT

Postpartum minor discomforts occur resulting from all systems adaptation. Perineal pain is one of the most common post partum minor discomforts. Cold therapy can help in reducing perineal pain after normal vaginal delivery. The main purpose of the study is to assess the effect of cold therapy for reduction of perineal pain among postnatal mothers in selected hospitals, Assam. With an evaluative approach, a pre-experimental, one group pretest-post-test design was adopted for the study among 60 samples by using purposive sampling technique. The study was conducted in Swahid Tilak Hemram Gunabhiram Civil Hospital, Morigaon and Bhurbandha Model Hospital, Morigaon, Assam. After obtaining consent from participants, pain score was assessed before application of cold therapy and it was applied for 15-20 min twice a day i.e immediately within 1 hour of normal vaginal delivery and subsequently after 8 hours for two consecutive days. The level of pain was assessed each day by using Numerical Pain Rating Scale. The study revealed that the mean postest level of pain (1.48) was lower than the mean pretest level of pain (7.73). The calculated ts=39.36 and p-value=0.001 was significant at p<0.05, thus research hypothesis (H₁) was accepted. The result of Chi square test and Fisher's Exact test revealed that age is associated with the pretest level of perineal pain. The present study concluded that cold therapy application was proved to be significant in reducing perineal pain after normal vaginal delivery among postnatal mothers.

Keywords: Perineal pain, episiotomy wound, perineal tear, cold therapy, numerical pain rating scale





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INTRODUCTION

Delivering a baby out of a relatively small opening can leave the vaginal and perineal muscle torn, with tears and sometimes the perineum is required to be given episiotomy and stitched. This causes a lot of perineal pain and discomfort [1]. Normal puerperium is 6 weeks post childbirth during which the pelvic organs reverts back to the near pre-pregnant state and the physiological changes of pregnancy are reversed [2]. WHO describes the postnatal period as most critical phase and yet most neglected period for the provision of quality care in the lives of mothers and babies; most death occur in the postnatal period [3]. Postpartum minor discomforts may occur resulting from all systems adaptation. Most common post partum minor discomforts include after pain, perineal pain, constipation, urinary distension, lactation problems. Immediate and effective care for these problems during intranatal and postnatal period can make the differences in postpartum adaptation [4]. It is the priority of the health care professionals to identify and value the morbidities resulting from a normal vaginal delivery, particularly the presence of pain. The identification of spontaneous perineal pain and the methods used for its relief are considered very important [5]. It is important that health care professional who attend to new postnatal mothers, should know how to assess and treat perineal pain. Considering the high rates of perineal trauma following vaginal deliveries, we as a midwife should need to provide cost-effective alternative treatment for perineal pain, based on scientific evidence. The researcher had been exposed to various hospitals and clinical settings where it was observed that women undergoing normal vaginal delivery with or without episiotomy had various degree of perineal pain and due to the pain the postnatal mothers find it difficult to do their daily activities of living and it interferes with caring the baby. The postnatal mothers, because of fear of pain do less voiding and passing of stool which leads to other issues during her stay in hospital. Non pharmacological treatment such as icepack covered with a soft covering can be used to reduce swelling and discomfort [6]. Cold therapy application is a very cost effective procedure and studies have shown the positive effect in relieving the discomfort of the perineal pain of the postnatal mothers. Moreover, this type of study has been done very less in the selected population. Therefore, the researcher felt the need to provide cold therapy application for perineal pain reduction after normal vaginal delivery among the postnatal mothers.

OBJECTIVES

- 1. To assess the level of perineal pain before cold therapy application among postnatal mothers in selected hospitals
- 2. To assess the level of perineal pain after cold therapy application among postnatal mothers in selected hospitals.
- 3. To assess the effect of cold therapy for reduction of perineal pain among postnatal mothers in selected hospitals.
- 4. To find out the association between level of perineal pain before cold therapy application among postnatal mothers with the selected demographic variables.

MATERIALS AND METHODS

A pre experimental one group pre-test-post-test design was selected for the study. The study was conducted in Swahid Tilak Hemram Gunabhiram Civil Hospital, Morigaon and Bhurbandha Model Hospital Bhurbandha, Morigaon among 60 postnatal mothers. Ethical clearance certificate and formal permission was taken from the concerned authorities and participants to conduct the research study. The tools used for the study was demographic variables and Mc Caffery Numerical Pain Rating Scale. Non-probability Purposive sampling technique was used for selecting the postnatal mothers who had normal vaginal delivery with episiotomy or perineal tear. The pre-test was conducted before cold therapy application by using Mc Caffery Numeric Pain Rating scale and cold therapy was applied for 15-20 min twice a day i.e immediately within 1 hour of normal vaginal delivery and subsequently after 8 hours for two consecutive days. Observation was done before and after the intervention and postest pain score was





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obtained once again for each selected samples. The data obtained were analyzed in terms of objective of the study by using descriptive and inferential statistics.

Inclusion criteria

The study includes those postnatal mothers

- ➤ Who are on the 1st postnatal day after normal vaginal delivery with episiotomy or perineal tear admitted in selected hospitals, Assam.
- ➤ Who are available during the time of data collection period.
- \triangleright Who are having pain score of ≥ 4 in numerical pain rating scale.
- ➤ Who are able to communicate in English, Assamese.

Exclusion criteria

The study excludes the postnatal mothers

- ➤ Who are not willing to participate in the study.
- > Who are receiving analgesic drug.
- > Who are using any traditional method for reduction of perineal pain.
- ➤ Who are with any kind of mental health or medical related issues.

RESULT

With reference to the sample characteristics presented in Table 1, most of the postnatal mothers were in the age group of 21-25 years, i.e, 40 (66%), 30(50%) had higher secondary level of education, 35 (58%) were primi gravida, 27 (45%) of the postnatal mothers were from rural area, 35 (58%) had episiotomy wound, 46 (76%) birth weight of newborn was in between 2.5-3 kg. With reference to the level of perineal pain of the postnatal mothers pretest and posttest perineal pain score result indicates that 92% had pain score of 7-10 (severe pain). Whereas in the posttest 82% had pain score of 1-3 (mild pain). Hence posttest pain score remains lower than the pretest pain score. With reference to the effect of cold therapy for reduction of perineal pain after normal vaginal delivery among postnatal mothers presented in Table 2, the results shows that the mean postest pain score (1.48) was lower than the mean pretest pain score (7.73). The calculated 15% = 39.36 and p-value = 0.001 was significant at p<0.05. Therefore, cold therapy application was proved to significantly reduce the perineal pain among postnatal mothers.

With reference to the association between the pretest level of perineal pain before cold therapy application among postnatal mothers with the selected demographic variables. The data indicates that there is a significant association between the pretest level of perineal pain with the age (in years). Hence, the research hypothesis H_2 is accepted and null hypothesis H_{02} is rejected.

DISCUSSION

The findings of the present study had been discussed in relation with the objectives and hypotheses of the study under following headings;

To assess the of level of perineal pain before cold therapy application among postnatal mothers in selected hospitals. The findings of the present study revealed that out of 60 postnatal mother: majority 92% (55) had pain score between 7-10 (severe pain) and 8 % (5) had pain score between 4-6 (moderate pain) before the application of cold therapy. Similar study conducted by Sadashivsali A (2013) on effect of sitz bath versus gugguldhupan on pain related to episiotomy in postnatal mothers admitted in selected hospitals of Pune city, India. The sample size was 60 postnatal mothers, divided into two groups and convenience sampling technique was used. Group A consisted of 30 samples and received sitz bath as a treatment. Group B had another 30 samples who received gugguldhupan as a treatment. The study revealed that, 53.3 % postnatal mothers having severe episiotomy pain before giving sitz bath and





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according to self assessment score maximum 63.3 % postnatal mothers having severe episiotomy pain before gugguldhupan [7].

To assess the of level of perineal pain after cold therapy application among postnatal mothers in selected hospitals

The present analysis revealed that out of 60 postnatal mother: majority 82% (49) had pain score between 1-3 (mild pain) and 18% (11) had 0 (no pain) after the application of cold therapy. Similar study conducted by Bini, Solomon JR, Ahita V (2019) on effectiveness of cold application on episiotomy pain among postnatal mothers in Kanyakumari district. The result revealed that experimental group in day 1 had pretest score of 53.33% very severe episiotomy pain and 46.67% worse episiotomy pain and on postest 73.34% severe pain and 13.33% very severe episiotomy pain. On day 2 the pretest pain score was 53.33% moderate pain and 46.67% severe pain which was reduced to 66.67% mild pain and 33.33% no episiotomy pain in postest. Whereas in the control group in day 1 pretest 53.33% very severe episiotomy pain and 46.67% worse episiotomy pain and on day 1 postest 90% had very severe pain, 6.67% had severe pain, 3.33% had worse episiotomy pain. On day 2 in pretest score 66.67% very severe pain and 33.33% severe pain and on postest 66.67 severe pain, 16.67% very severe pain, 16.67% moderate pain. There was a significant reduction in mean post-test episiotomy pain of the experimental group [8].

To assess effect of cold therapy for reduction of perineal pain among the postnatal mothers in selected hospitals.

Finding of the present study reveals that the mean postest level of pain (1.48) was lower than the mean pretest level of pain (7.73). The calculated t_{59} value was 39.36 and the tabulated valued of df for 0.05 significance was 0.001 was significant at p<0.05, thus the null hypothesis (H₀₁) was rejected and research hypothesis (H₁) was accepted. Therefore, cold therapy application was proved to significantly reduce the perineal pain after normal vaginal delivery among postnatal mothers.

Similar study conducted by Senol KD, Aslan E (2017) to determine the effects of cold application to the perineum on pain relief after vaginal birth in Turkey among 200 postnatal mothers. The samples were divided into experimental group and control group. Experimental group got application of crushed ice pack for 20 min in the postpartum 1st 2 hours and 4 hours after the first application. The result showed that the experimental group, the first visual analog scale score was 6.7±1.68; after cold gel pad application, the pain level decreased to 2.59±1.20 in both primiparous and multiparous mother. The result showed that both episiotomy and perineal laceration were strongly associated with the presence of perineal pain during immediate postpartum and at 3 months. It also showed that increased length of labor and decrease parity have been positively with postpartum perineal [9].

To find out the association between level of perineal pain before cold therapy application among postnatal mothers with the selected demographic variables.

The association findings of the present study revealed that there was a significant association between the pretest level of pain of pain with the age (in years). Hence, the research hypothesis H_2 is accepted and null hypothesis H_0 is rejected. Similar study conducted by Mathias ADREA, Pitangui RCA, Vasconcelos AMA, Silva SS (2015) on perineal pain measurement in the immediate vaginal postpartum period in Petrolina and Juazerio among 147 women who had delivered baby through vaginal delivery. McGill Pain Questionnaire (MPQ) and numeric visual pain scale (NVS) was used to measure pain. The data was collected at a minimum of 6 hours and maximum of 24 hours after delivery. The study revealed that there was a significant association of pain with episiotomy (p=0.002), laceration (0.039) and parity (0.028). Thus the study concluded that there was significant association of pain with episiotomy, laceration, parity [10].





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CONCLUSION

The present study was conducted to evaluate the effect of cold therapy for reduction of perineal pain among postnatal mothers. The findings of the study revealed that cold therapy is effective for reduction of perineal pain among postnatal mothers.

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Table 1: Frequency and percentage distribution of demographic variables

n=60

Sl. No.	Demographic Variables	Frequency (f)	Percentage (%)
1	Age (in years)		
	a) <u><</u> 20	11	18
	b) 21-25	40	67
	c) ≥26	9	15
2	Educational status		
	a) Primary	-	-
	b) High school	17	28
	c) Higher secondary	30	50
	d) Graduation and above	13	22
		-	-
2	e) No formal education		
3	Gravida		
	a) Primi gravida	35	58
	b) Multi gravida	25	42
4	Area of residence		





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	a) Urban	20	33	
	b) Rural	27	45	
	c) Semiurban	13	22	
5	Type of perineal wound			
	a) Perineal tear wound	25	42	
	b) Episiotomy wound	35	58	
6	Birth weight of newborn (in kg)			
	a) ≤2.4`	-	-	
	b) 2.5 – 3.0	46	77	
	c) 3.1 - ≥3.5	14	23	

Table 2: Frequency and percentage distribution of pretest and postest level of perineal pain among postnatal mothers

n=60

Level of perineal pain	Pre-test		Post-test	
	F	%	f	%
No pain	-	-	11	18
Mild pain	-	-	49	82
Moderate pain	5	8	-	-
Severe pain	55	92	-	-
Total	60	100	60	100

Table 3: Mean, Standard Deviation, Mean Difference, t test value, df, p value of pain score before and after cold therapy application among postnatal mothers.

n=60

Comparison of level of pain	Mean	SD	Mean Difference	t value	df	p value	Inferences
Pre-test	7.73	1.006					
Post-test	1.48	0.965	6.25	39.36	59	0.001*	S

^{*} Significant at p<0.05 (t59=39.36)

S-Significant

Table 4: Association between level of perineal pain before cold therapy application among postnatal mothers with the selected demographic variables

n=60

Sl.	Demographic variables	Pre-test level of perineal pain		χ²	df	p value Fisher's	Inference
No	0 1	Moderate	Severe	Value		exact test	
	Age (in years)						
1	a. ≤20	3	8	3.653	1	0.038*	S
	b. ≥21	2	47	3.633	1		3
	Educational status						
2	a. High school	1	16			1.00	
	b. Higher secondary and	4	39	0.186	1	1.00	NS
	above						





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3	Gravida a. Primi gravida	3	32	0.006	1	1.00	NS
4	b. Multi gravida Area of residence a. Urban	1	23 19			0.656	
	b. Others	4	36	0.436	1		NS
5	Type of perineal wound a. Perineal tear wound	2	23	0.006	1	1.00	NS
	b. Episiotomy wound	3	32	0.000	1		110
	Birth weight of newborn (in kg)						
6	a. ≤3.0 b. ≥3.1	4 1	42 13	0.034	1	1.00	NS

^{*} Significant at p<0.05



S-Significant NS-Non Significant



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RESEARCH ARTICLE

Customer Perception towards Township Apartment

A. Pappu Rajan^{1*}, P.J. Rosario Vasantha Kumar² and Sunil Raj Y³

¹Associate Professor, Department of Management Studies, St. Joseph's Institute of Management, St. Joseph's College (Autonomous), (Affiliated to Bharathidasan University), Tiruchirappalli, Tamil Nadu,

²Librarian, St. Xavier's College (Autonomous) Palayamkottai, (Affiliated to Manonmaniam Sundaranar University), Tirunelveli, Tamil Nadu, India.

³Assistant Professor, Department of Data Science, St. Xavier's College (Autonomous) Palayamkottai, (Affiliated to Manonmaniam Sundaranar University), Tirunelveli, Tamil Nadu, India.

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*Address for Correspondence

A. Pappu Rajan

Associate Professor,

Department of Management Studies,

St. Joseph's Institute of Management, St. Joseph's College (Autonomous),

(Affiliated to Bharathidasan University),

Tiruchirappalli, Tamil Nadu, India.

Email: ap_rajan2001@jim.ac.in



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ABSTRACT

Housing plays a crucial role in the economic, social, and civic development of both developed and developing countries. This study delves into the perceptions of middle-class individuals regarding living in apartment townships, particularly focusing on their understanding of affordable housing provided by developers in Chennai, India. By directly engaging with residents, the research aims to gain insights into their attitudes towards affordable housing options. The study specifically targets middle-income individuals who have had experience with apartment living in Chennai. Employing a descriptive research design, it seeks to analyze the market dynamics, enhance public comprehension of affordable housing concepts, and potentially generate more leads for banks. Objectives include assessing market demand, identifying potential risks for banks, and proposing mitigation strategies. Data collection involves simple random sampling from a database of builder projects in Chennai. Analysis is conducted using Microsoft Excel and IBM SPSS software, employing statistical measures such as means, descriptive statistics, and variance analysis to interpret primary data sources accurately.

Keywords: Segmentation, Customer perception, Marketing Research, Consumer Behavior





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INTRODUCTION

A mediation model (Yingzi Xu et.al,2006) is used to links customer perceived service value to customer loyalty via customer satisfaction. The customer satisfaction (Boamah, Fredrick Ahenkora,2020) is impact of different combination of reliability, responsiveness, assurance and empathy with high service quality. Market Segmentation includes psychographics, lifestyles, values, behaviours, and multivariate cluster analysis routines of individual perceptions and experiences. According to Bharwana(2013), service quality is very high impact on customers' satisfaction. Market segmentation Process has many difficulties that to be solved and it has described to be tackled to leading several management teams (Dibb, S. and Simkin, L,1997) through the analysis and strategy. As a result of the ongoing economic deregulation measures of the Government particularly those in the financial and banking sector, there have been market changes in the housing finance systems in India (P K, Manoj,2004). The House (Mondal et.al. 2020, Balanaga Gurunathan K and Nidhi Ahuja (2020) is the key need that the livelihood of a person as a proper house gives an individual security of the people. In this research gives over all view of customer perception towards township apartments and try to deal the important of the research in the filed of housing finance in particular to township apartments.

Conceptual Frame Work

Housing stands as a vital catalyst for economic, social, and civic development, constituting a significant component of developing nations' economies. The provision of public transport, parks, and local amenities is crucial for fostering community well-being within apartment complexes. While luxury amenities are valued, practicality and quality in personal and shared living spaces take precedence for most individuals (Hanif, 2015). Intense competition within the banking sector necessitates unwavering commitment to service excellence (Bose Ranjan, 2018). Urban density fluctuations and transportation demand dynamics influence housing preferences and purchasing behavior (Habibia& Asadi, 2011). Affordability has emerged as a central theme in housing policy and urban sustainability discussions (Mubiru &Ikiriza, 2021; Gupta, 2019). Economic models often focus on factors like transport accessibility and house prices in understanding housing markets (Matenge et al., 2016). High-rise living impacts various aspects of residents' lives, tempered by socioeconomic factors and neighborhood quality (Gifford, 2007). Location remains pivotal in the housing market, with proximity to employment and amenities driving demand (Aluko, 2011). The quality of economic output serves as a critical indicator for strategic business planning and consumer behavior (Ilieska, 2013). Understanding customer perceptions guides real estate developers in meeting buyer demands effectively (Misra et al., 2013; Engel et al., 2005). Open spaces within housing societies enhance residents' quality of life and property values (Kakkar & Supriya, 2014). Factors such as landownership, zoning, and public utilities are key considerations in apartment site selection for developers (Krisnaputri, 2016). As cities globally brace to accommodate an additional 3 billion people over the next two decades, the lack of resources poses a significant challenge to sustainable urban development.

Research Design

The study is focused on understanding the perceptions of middle-class individuals towards apartment township living and their comprehension of affordable housing provided by developers. It aims to delve into the depth of general awareness of affordable housing concepts prevalent in the market and evaluate the extent of market penetration achieved by builders. Conducted in Chennai, India, the research targets middle-income residents who have direct experience with apartment living. By engaging with this demographic, the study seeks to unravel the relationship between their demographic profiles and their perceptions of affordable housing.

The methodology employed incorporates a descriptive research design, allowing for a comprehensive analysis of various factors influencing perceptions, including physical, social, and economic attributes associated with apartment living. Additionally, the study endeavors to identify the preferences and requirements of respondents concerning ideal apartment living arrangements. Through this process, the research aims to contribute to a deeper understanding of the affordable housing market dynamics in Chennai.





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The primary objective of the study is to provide insights into the affordable housing market, thereby facilitating enhanced understanding among both consumers and stakeholders, potentially leading to increased leads for HFFC (Housing Finance and Development Corporation). Apart from this overarching goal, the study also seeks to achieve several specific objectives. These include analyzing current market trends, assessing demand and supply dynamics, examining pricing mechanisms, and identifying potential risks faced by HFFC in the market. Furthermore, the research aims to propose mitigation strategies to address these risks effectively. To accomplish these objectives, the study employs a systematic approach to sample selection and data collection. The target population comprises individuals aged 20 to 60 years, belonging to the middle-income bracket, with monthly household incomes ranging from Rs. 5000 to Rs. 30000. These individuals must have been residents of Chennai and its surrounding areas for at least two years. The sampling technique employed is simple random sampling, drawing data from a comprehensive database of builder projects in Chennai. This ensures that the sample is representative of the broader population of interest. Data collection involves both primary and secondary sources. Primary data is collected through direct field visits and personal interviews using structured questionnaires. These interviews are conducted with randomly selected respondents residing in areas covered by builder projects listed in the database.

On the other hand, secondary data is gathered from HFFC's database, focusing on builder projects in specific areas such as Avadi and Ambattur, which cater to the affordable housing segment. The collected data is then subjected to rigorous analysis using appropriate statistical tools and software. Microsoft Excel is utilized for data input and initial organization, while IBM SPSS (Statistical Package for the Social Sciences) is employed for in-depth statistical analysis. Various statistical techniques are applied, including means comparison, descriptive statistics, variance analysis, and Chi-Square tests for goodness of fit. These analytical methods help in uncovering patterns, trends, and associations within the data, providing valuable insights into the perceptions and preferences of respondents. One of the critical aspects of the study is the examination of the relationship between respondents' demographic profiles and their perceptions of affordable housing. This involves testing the association between different demographic variables and the dependent variable of perception. By analyzing this relationship, the study aims to identify any significant demographic factors that influence individuals' perceptions of affordable housing. Furthermore, the research explores the concept of market potential in the affordable housing segment in Chennai.

It assesses the demand for affordable housing, considering factors such as population growth, income levels, and urbanization trends. Additionally, the study examines supply-side factors, including the availability of land, construction costs, and regulatory frameworks governing the housing sector. By analyzing both demand and supply dynamics, the research aims to provide a comprehensive understanding of the market landscape. Another important aspect of the study is the assessment of potential risks faced by HFFC in the affordable housing market. This involves identifying various risk factors, such as economic downturns, changes in government policies, and market competition. Subsequently, the research proposes mitigation strategies to manage these risks effectively and safeguard the interests of HFFC and other stakeholders. In summary, the study aims to contribute to a deeper understanding of the affordable housing market in Chennai by examining middle-class perceptions and preferences. Through a systematic research approach encompassing data collection, analysis, and interpretation, the study seeks to provide valuable insights for consumers, developers, policymakers, and financial institutions operating in the housing sector. By shedding light on market dynamics, potential risks, and mitigation strategies, the research aims to foster informed decision-making and promote sustainable growth in the affordable housing segment.

RESEARCH DISCUSSION

The researcher is tried to find the reason for choosing apartment as result of finding table 1.1 shows that 13 of respondents feel it is affordable, 45 of respondents feel it is proximity to transport and other amenities, 15 of respondents feel it is a socialistic thought who have responded for the survey. The researcher is tried to find the reason for shift from apartment as result of finding table 1.2 shows that 24 of the respondents shift due to rent or maintenance problem, 27 of the respondents shift due to independent living, 19 of the respondents shift due to





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Quality issues and 14 of the respondents shift due to other reasons who have responded for the survey. Figure 1, presents the details on the reasons for the respondents to move form the apartment to another. The peak shows that most respondents would leave from the apartment due to the quality issues. H1:There is no significant difference between the perception of the people towards affordable apartment township living and their gender. As the P value is 0.094 which is lesser than 0.1 it shows that gender has influence on perception towards township apartments. Table 1.4 presents symmetric measures examining the relationship between gender and perception towards township living. Pearson's correlation coefficient for interval by interval variables is 0.034, while Spearman's rank correlation coefficient for ordinal by ordinal variables is 0.036. Both coefficients suggest a weak correlation between gender and perception towards township living. The asymptotic standard errors for both coefficients are 0.100. The approximate t-values are 0.335 for Pearson's correlation and 0.354 for Spearman's correlation. The p-values for both coefficients are notably high, with 0.738 for Pearson's correlation and 0.724 for Spearman's correlation, indicating that the observed correlations are not statistically significant at the 5% significance level. Thus, based on this analysis, there is insufficient evidence to support a significant relationship between gender and perception towards township living in the dataset. Figure 2, presents the symmetric measures where gender and perception towards township are considered. The peak shows the significance of the results, standard error and the values associated.

The Pearson's correlation coefficient of 0.034 indicates a positive relationship between gender and perception towards township living. Multiple hypotheses were tested, revealing that: a) educational qualifications have minimal impact on perception towards township apartments, b) salary influences perception towards such apartments, and c) employment type also affects perception. The questionnaire developed to assess customer attitudes towards apartment townships in the affordable housing segment proved valuable in gauging customer awareness and perception of available products. It can be utilized for future studies and market analyses to further understand individual perceptions. The sample population skewed male, with 61% male respondents compared to 39% female, reflecting the predominant male presence at HFFC branches. Age emerged as a significant determinant of perception, with those aged 20-30 exhibiting higher satisfaction with apartment townships than those aged 50-60. Educational qualifications, job type, and income were found to be interrelated factors influencing customer perception, with individuals of higher qualifications, income, and job status expressing less satisfaction with apartment townships. Conversely, those with moderate educational backgrounds, middle-class incomes, and decent jobs showed a more positive outlook. Among these factors, the locality and price of apartment townships exerted the most significant impact. Private sector banks should focus on enhancing customer satisfaction by offering housing loans with simplified procedures and transparent terms. Diversifying product offerings and leveraging various media channels for promotion is essential. Understanding and addressing customer grievances regarding home loans, including high-interest rates, through interest subsidies based on income categories, would benefit both customers and the organization. Banks should closely monitor customer preferences and collaborate with builders, DSAs, and masons to stay abreast of customer needs and generate more leads. Continuous promotion of products, including sponsoring builder community meetings, is vital for maintaining visibility and market presence.

CONCLUSION

The housing sector is an ever-expanding industry, heavily reliant on meeting customer needs and satisfaction for success. Any decrease in customer satisfaction and comfort within this sector could result in significant losses. Apartment townships have emerged as one of the fastest-growing segments within India's housing market, playing a pivotal role in revenue generation and contributing substantially to the country's GDP. Despite the industry's apparent success and sustained growth, there exists a notable challenge stemming from individual perceptions and viewpoints, which influence family members' attitudes towards the apartment township lifestyle. While younger demographics, such as teenagers, those in their early twenties, and thirties, often express a desire to reside in apartment townships, individuals aged forty and above typically exhibit strong reluctance toward embracing this lifestyle. Furthermore, there's a noticeable divide based on socioeconomic status and education level, with affluent and highly educated individuals displaying hesitancy while middle-class individuals with moderate educational





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backgrounds show greater interest, relatively speaking. Our research has identified several key factors that significantly impact customers' inclination towards apartment townships. Privacy emerges as a primary concern, as residents may not enjoy the same level of privacy as they would in standalone houses. Despite the affordability of apartment townships, peace and proximity to amenities also play essential roles in shaping people's perceptions of this housing option. Therefore, it is imperative to focus efforts on addressing these concerns and improving customer satisfaction while working to alter perceptions of apartment townships.

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Table 1 Reason for choosing apartment

		Frequency	Percent	Valid Percent	Cumulative Percent
	Affordable	13	13.0	13.0	13.0
	Proximity to transport and other amenities	45	45.0	45.0	58.0
Valid	lifestyle and city living, better safety and security	15	15.0	15.0	73.0
	socialistic thought	27	27.0	27.0	100.0
	Total	100	100.0	100.0	

Table 2 Reason for shift from apartment

		Frequency	Percent	Valid Percent	Cumulative Percent
	Rent or maintance problem	24	24.0	24.0	24.0
Valid	wanted independent living	27	27.0	27.0	51.0
	Proximity problems	19	19.0	19.0	70.0





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Quality Issues	16	16.0	16.0	86.0
others	14	14.0	14.0	100.0
Total	100	100.0	100.0	

Table 3. Relationship between gender and Perception towards township apartments

			Asymptotic
	Value	Df	Significant (2 Sided)
Pearson Chi-Square	1.242a	4	.094
Likelihood Ratio	1.194	5	.094
Linear-by-Linear Association	.113	1	.073
N of Valid Cases	100		

Table 4 Symmetric Measures – Gender and perception towards township

	Value			Approximate	Approximate
value			Error	T^b	Significance
Interval by	Pearson's R	.034	100	.335	.738°
Interval	rearson's K	.034	.100	.333	./30°
Ordinal by	Spearman	.036	.100	.354	.724°
Ordinal	Correction	.036	.100	.554	./24
N of Valid Cases 100					

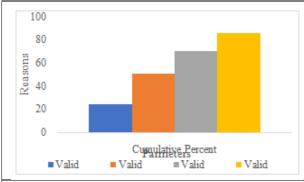


Figure 1. Reason for Shift between apartment

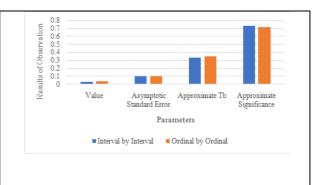


Figure 2. Symmetric Measures – Gender and perception towards township





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RESEARCH ARTICLE

For Multi-Valued Mappings, Fixed Point Theorems in Complex Valued **b-Metric Spaces**

Arul Xavier A* and Maria Joseph J

Department of Mathematics, St.Joseph's College, (Affiliated to Bharathidasan University Tiruchirappalli). Tiruchirappalli, Tamil Nadu, India.

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*Address for Correspondence Arul Xavier A

Department of Mathematics,

St.Joseph's College,

(Affiliated to Bharathidasan University Tiruchirappalli).

Tiruchirappalli, Tamil Nadu, India.

Email: arulxavier3006@gmail.com,



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ABSTRACT

This study establishes the complex-valued b-metric spaces on the fixed point theorem for two multivalued mappings, which generalizes several papers in the present literature.

Keywords: Fixed point, contraction, complex-valued metric space, set-valued mapping, complete metric space, b-metric space.

INTRODUCTION

In mathematical analysis, the fixed point theory is crucial. In 1922, Stefan Banach[1] proposed contraction mapping principle which established foundation for many results fixed point theory. Subsequently, many researchers[2], [3][4]generalized the principle of Banach contracture mapping in metric and b-metric spaces. Regards, multi-valued transformations in b-metric spaces, Josephet.all[5] proved the fixed point theorem. Then Saleh AAl-Mezelet.all[6] proved fixed point, common fixed point theorem on multi-valued mappings in complex-valued metric spaces. This article, explores invariant point, common fixed point theorems on multiple valued transformations in complexvalued b-metric spaces.

Definition 1.1.Consider $\mathbb C$ stands for the collection of numbers in complex planebesides z_1 , z_2 ϵ $\mathbb C$. On $\mathbb C$, consider a partial order ≤ as follows:

 $z_1 \le z_2$ iff $Re(z_1) \le Re(z_2)$ and $Im(z_1) \le Im(z_2)$. Thus $z_1 \le z_2$ if one of the following holds:

- 1. $Re(z_1) = Re(z_2)$ and $Im(z_1) = Im(z_2)$;
- 2. $Re(z_1) < Re(z_2)$ and $Im(z_1) = Im(z_2)$;





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- 3. $Re(z_1) = Re(z_2)$ and $Im(z_1) < Im(z_2)$;
- 4. $Re(z_1) < Re(z_2)$ and $Im(z_1) < Im(z_2)$;

We will write $z_1 \lesssim z_2$ if $z_1 \neq z_2$ and one of (2), (3), and (4) is satisfied; also we'll write $z_1 \prec z_2$ only if (4) is met. It shows that

- i. $0 \le z_1 \le z_2$ implies $|z_1| < |z_2|$;
- ii. $z_1 \le z_2$ as well as $z_2 < z_3$ imply $z_1 < z_3$;
- iii. $0 \le z_1 \le z_2$ implies $|z_1| < |z_2|$;
- iv. if $a, b \in \mathbb{R}$, $0 \le a \le b$ and $z_1 \le z_2$ then $az_1 < bz_2$ for all z_1 , $z_2 \in \mathbb{C}$.

Definition1.2.Assume *X* be a set of non-empty and $d: X \times X \to \mathbb{C}$ is known as a complex metric on *X*, if for every $x, y, z \in X$, the conditions holds are

- i. $0 \le d(x, y)$ with d(x, y) = 0 iff x = y,
- ii. d(x, y) = d(y, x),
- iii. $d(x,y) \le d(x,z) + d(z,y)$.

The pair (X,d) is then referred to a metric space in complex-valued .

Definition 1.3.Assuming $\{x_n\}$ be a sequence in X and $x \in X$. On every $c \in \mathbb{C}$, and 0 < e there is $n_0 \in \mathbb{R}$ in order toall $n \ge n_0$, $d(x_n, x) < c$, then assume that x_n converges to x. If $x_n \to x$, then we call x is the boundary point of x_n . We denotes limit by $\lim x_n = x$ or $x_n \to x$ when **Definition 1.4.**Let $\{x_n\}$ be a sequence in X. If for every $c \in \mathbb{C}$, with 0 < c there is $n_0 \in \mathbb{N}$ such that for all $n \ge n_0$, $d(x_n, x_{n+m}) < c$, then x_n is called Cauchysequence in (X, d). Every sequence of Cauchy in entiremetric space (X, d) is convergent.

Definition 1.5. [1] Letting(X, d) be a metric space with complex-values and asequence x_n belongs to X. Therefore x_n converges to x iff $|d(x_n, x)| \to 0$ as $n \to \infty$.

Lemma 1.6. [1] Consider(X, d) be a metric space with complex-valued also x_n be asequence in X. Thus x_n Cauchy sequence iff $|d(x_n, x_{n+m})| \to 0$ as $n \to \infty$.

Definition1.7. $\mathcal{H}: \mathcal{CB}(X) \times \mathcal{CB}(X) \to [0, \infty)$ is a function derived by $\mathcal{H}(A, B) = \max\{\sup_{x \in A} D(x, A), \sup_{x \in B} D(x, B)\}$ is the Hausdorff metric on $\mathcal{CB}(X)$ caused by the metric d on X here $D(x, A) = \inf\{d(x, y) : y \in A\}$ for every $A \in \mathcal{CB}(X)$.

Definition 1.8. If v = Tv (when $T : X \to X$ is a single-valued map) or $v \in Tv$ (while $T : X \to \mathcal{P}(X)$ is a map with several values), there is av inX could be a fixed point of a map T. We states T has an endpoint if there contains $v \in X$ such that $Tv = \{v\}$. The collection of invariant points of T is represents T and the group of two multi-valued mappings, common fixed points T, T0 are denotes T1.

Definition 1.9. Two mappings with multiple values $T, S: X \to \mathcal{CB}(X)$, claims that T, S fulfil the property of common approximate endpoint if there is a $\{x_n\} \subset X$ in the sense that $\lim_{n \to \infty} \mathcal{H}(\{x_n\}, Tx_n) = \lim_{n \to \infty} \mathcal{H}(\{x_n\}, Sx_n) = 0$

Definition 1.10.For two mappings T, S: $X \to X$, we say that T, Sshare a common arbitrary fixed point if there consists a sequence $\{x_n\} \subset X$ such that

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\lim_{n\to\infty} d(\lbrace x_n\rbrace, Tx_n) = \lim_{n\to\infty} d(\lbrace x_n\rbrace, Sx_n) = 0
```

Throughout the paper, we assume that $\{a, b, c, d, e\} \subset [0,1)$. This article's fundamental theorem is as follows.

Theorem 1.11.Assume that (X, d) be an entire metric space besides $T, S: X \to \mathcal{CB}(X)$ are two multi-valued functions in order to every $x, y \in X$.





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 $\mathcal{H}(Tx, Ty) \leq$

$$ad(x,y) + bD(x,Sx)D(y,Ty) + c\sqrt{D(y,Sx)D(x,Ty)} + dD(x,Sx)D(x,Ty) + eD(y,Sx)D(y,Ty).$$

where a + b + c + 2d + 2e < 1. If and only if T, Scan satisfies the common approximate end point feature, hence they have a unique end point.

Theorem 1.12.Assuming an absolute metric space(X, d)alsoT, $S: X \to \mathcal{CB}(X)$ be two multiple valued functions in the sense that each x, $y \in X$.

$$\mathcal{H}(Tx,Ty) \leq ad(x,y) + b \frac{D(x,Sx)D(y,Ty)}{1+d(x,y)} + c \frac{\sqrt{D(y,Sx)D(x,Ty)}}{1+d(x,y)} + d \frac{D(x,Sx)D(x,Ty)}{1+d(x,y)} + e \frac{D(y,Sx)D(y,Ty)}{1+d(x,y)}$$

where a + b + c + 2d + 2e < 1. Then T, S having an endpoint uniquely, iffthey fulfils the common property of an approximate endpoint.

Now, we can present our key findings.

PRIMARY FINDINGS

Theorem 2.1.Let (W, d) be a entire valued space of b-metric with $s \ge 1$ also assume that $F, G: W \to \mathcal{CB}(W)$ be two multiple valued functions in the sense that toevery $a, b \in W$.

 $d(Fb,Ga) \le k_1 d(a,b) + k_2 d(a,Ga)d(b,Fb) + k_3 d(b,Ga)d(a,Fb) + k_4 sd(a,Ga) + d(a,Fb) + k_5 d(b,Ga)d(b,Fb)$ where $k_1 + k_2 + k_3 + 2k_4 s + 2k_5 < 1$. $k_i \ge 0$, i = 1,2,3,4,5. Thus F, G having a fixed point uniquely.

Proof. Letting $a_0 \in W$ be a random element and also $a_{2n-1} = Fa_{2n-2}$. We have

$$d(a_{2n+1}, a_{2n}) = d(Fa_{2n}, Ga_{2n-1})$$

$$\leq k_1 \, d(a_{2n} \, , a_{2n-1}) + \, k_2 \, d(a_{2n-1} \, , Ga_{2n-1}) d(a_{2n} \, , Fa_{2n}) + \, k_3 d(a_{2n} \, , Ga_{2n-1}) d(a_{2n-1} \, , Fa_{2n}) \\ + \, k_4 \, s d(a_{2n-1} \, , Ga_{2n-1}) d(a_{2n-1} \, , Fa_{2n}) + \, k_5 \, d(a_{2n} \, , Ga_{2n-1}) d(a_{2n} \, , Fa_{2n})$$

$$=k_1\,d(a_{2n}\,,a_{2n-1})+k_2\,d(a_{2n-1}\,,a_{2n})d(a_{2n}\,,a_{2n+1})+k_3d(a_{2n}\,,a_{2n})d(a_{2n-1},a_{2n+1})\\+k_4\,sd(a_{2n-1}\,,a_{2n})d(a_{2n-1},a_{2n+1})+k_5\,d(a_{2n}\,,a_{2n})d(a_{2n}\,,a_{2n-1})$$

$$= k_1 d(a_{2n}, a_{2n-1}) + k_2 d(a_{2n-1}, Ga_{2n}) d(a_{2n}, a_{2n+1}) + k_4 s d(a_{2n-1}, a_{2n}) (d(a_{2n-1}, a_{2n+1}) + d(a_{2n}, a_{2n+1}))$$

$$\leq k_1 d(a_{2n}, a_{2n-1}) + k_2 d(a_{2n}, a_{2n+1}) + k_4 sd(a_{2n-1}, a_{2n})$$

 $(d(a_{2n-1}, a_{2n})d(a_{2n-1}, a_{2n+1}))$

$$\leq k_1 \, d(a_{2n} \, , a_{2n-1}) + k_2 \, d(a_{2n} \, , a_{2n+1}) + k_4 \, s(d(a_{2n-1} \, , a_{2n}) d(a_{2n-1}, a_{2n+1})) \\ d(a_{2n+1-} \, , a_{2n}) = \frac{k_1 + k_4 \, s}{1 - k_2 - k_4 \, s} \, d(a_{2n-1} \, , a_{2n})$$

$$d(a_{2n-1}, a_{2n}) = d(Fa_{2n-2}, Ga_{2n-1})$$

$$\leq k_1 d(a_{2n-2}, a_{2n-1}) + k_2 d(a_{2n-1}, Ga_{2n-1}) d(a_{2n-2}, Fa_{2n-2}) + k_3 d(a_{2n-2}, Ga_{2n-1}) d(a_{2n-1}, Fa_{2n-2}) + k_4 s d(a_{2n-1}, Ga_{2n-1}) d(a_{2n-1}, Fa_{2n-2}) + k_5 d(a_{2n-2}, Ga_{2n-1}) d(a_{2n-2}, Fa_{2n-2})$$

$$=k_1\,d(a_{2n-2}\,,a_{2n-1})+\,k_2\,d(a_{2n-1}\,,a_{2n})d(a_{2n-2}\,,a_{2n-1})+\,k_3d(a_{2n-2}\,,a_{2n})d(a_{2n-1},a_{2n-1})\\ +\,k_4\,sd(a_{2n-1}\,,a_{2n})d(a_{2n-1},a_{2n+1})+\,k_5\,d(a_{2n-2}\,,a_{2n})d(a_{2n-2},a_{2n-1})$$

$$= k_1 d(a_{2n-2}, a_{2n-1}) + k_2 d(a_{2n-1}, Ga_{2n}) d(a_{2n-2}, a_{2n-1}) + k_5 d(a_{2n-2}, a_{2n}) d(a_{2n-2}, a_{2n-1})$$

$$\leq k_1 d(a_{2n-2}, a_{2n-1}) + k_2 d(a_{2n_1}, a_{2n}) + k_5 sd(a_{2n-2}, a_{2n-1})$$

 $(d(a_{2n-1}, a_{2n})d(a_{2n-2}, a_{2n+1}))$





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$$\leq k_1 d(a_{2n}, a_{2n-1}) + k_2 d(a_{2n}, a_{2n}) + k_5 s(d(a_{2n-2}, a_{2n-1})d(a_{2n-1}, a_{2n}))$$

$$d(a_{2n-1}, a_{2n}) = \frac{k_1 + k_5 s}{1 - k_2 - k_5 s} d(a_{2n-2}, a_{2n_1})$$

Now taking,
$$\lambda = max\left\{\frac{k_1 + k_4 s}{1 - k_2 - k_4 s}, \frac{k_1 + k_5 s}{1 - k_2 - k_5 s}\right\} < 1$$

We conclude that for each $n \in \mathbb{N}$, $d(a_{2n+1}, a_{2n}) \le \lambda d(a_n, a_{n-1})$. By a conventional technique, one can demonstrate that $\{a_n\}$ is Cauchy sequence. Because (W, d) is an entiremetric space, consists of $u \in W$ in order to $a_n \to u$.

Claim:Fu and u are identical to each other.

$$d(u, Fu) = d(u, Ga_{2n-1}) + d(Ga_{2n-1}, Fu)$$

$$\leq d(u,Ga_{2n-1}) + k_1 d(Ga_{2n-1},u) + k_2 d(Ga_{2n-1},GGa_{2n-1})d(u,Fu) + k_3 d(u,a_{2n-1})d(a_{2n-1},Fu) \\ + k_4 sd(a_{2n-1},Ga_{2n-1})d(a_{2n-1},Fu) + k_5 d(u,Ga_{2n-1})d(u,Fu)$$

Assuming *n* → ∞ from the above inequality, we get $d(u, Fu) \le 0$.

Thus Fu = u.

Similarly, Gu = u can deduce. Thus, it concludes u be the shared fixed point of F as well as G.

Corollary 2.1. Assume that a complete valued metric space (W, d) also letting $F, G: X \to \mathcal{CB}(W)$ be two functions of multivalued such that for all $a, b \in W$.

$$d(Fb,Ga) \le k_1 d(a,b) + k_2 d(a,Ga)d(b,Fb) + k_4 sd(a,Ga) + d(a,Fb) + k_5 d(b,Ga)d(b,Fb)$$
 where $k_1 + k_2 + 2k_4 s + 2k_5 < 1$. $k_i \ge 0$, $i = 1,2,3,4,5$. Thus F , G contains an invariant point commonly.

Theorem 2.2.An entire valued metric space(W, d) besides $s \ge 1$ also F, $G: W \to W$ be two functions and $k_i \ge 0$, i = 1,2,3,4,5 be such that $k_1 + k_2 + k_3 + 2k_4 + k_5 < 1$.

$$1,2,3,4,5 \text{ be such that } k_1 + k_2 + k_3 + 2k_4 s + 2k_5 < 1.$$
 Let $d(Fb,Ga) \le k_1 d(a,b) + k_2 \frac{d(a,Ga)d(b,Fb)}{1+d(a,b)} + k_3 \frac{d(b,Ga)d(a,Fb)}{1+d(a,b)} + k_4 s \frac{d(a,Ga)+d(a,Fb)}{1+d(a,b)} + k_5 \frac{d(b,Ga)d(b,Fb)}{1+d(a,b)}$

for all $a,b \in W$. Then $Fand \ G$ share a common fixed point.

Proof. We have
$$k_1 + k_2 + k_3 + k_4 s + k_5 < k_1 + k_2 + k_3 + 2k_4 s + 2k_5 < 1$$
.
 $\leq k_1 d(a,b) + k_2 \frac{d(a,Ga)d(b,Fb)}{1+d(a,b)} + k_3 \frac{d(b,Ga)d(a,Fb)}{1+d(a,b)} + k_4 s \frac{d(a,Ga)+d(a,Fb)}{1+d(a,b)} + k_5 \frac{d(b,Ga)d(b,Fb)}{1+d(a,b)}$
 $\leq k_1 d(a,b) + k_2 d(a,Ga)d(b,Fb) + k_3 d(b,Ga)d(a,Fb) + k_4 s d(a,Ga)d(a,Fb) + k_5 d(b,Ga)d(b,Fb)$

By theorem 2.1 we declares that F & G shares a common fixed point. Trivially, uniqueness shown verbatim as per demonstration of corollary 2.1.

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RESEARCH ARTICLE

A Nonlinear Mathematical Model for Asthma: Exploring the Impact of **Environmental Pollution**

N. Vasuki1* and D. Jasmine2

¹Research Scholar, PG & Research Department of Mathematics, Bishop Heber College (Affiliated to Bharathidasan University), Trichy, Tamil Nadu, India.

²Assistant Professor, PG & Research Department of Mathematics, Bishop Heber College (Affiliated to Bharathidasan University), Trichy, Tamil Nadu, India.

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*Address for Correspondence

N. Vasuki

Research Scholar,

PG & Research

Department of Mathematics,

Bishop Heber College (Affiliated to Bharathidasan University),

Trichy, Tamil Nadu, India.

Email: vasu681999@gmail.com



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ABSTRACT

This paper utilizes a nonlinear mathematical model to examine the effects of cigarette smoking and environmental pollution on the prevalence of asthma. To comprehend this relationship, we obtain an approximate analytical solution using the homotopy perturbation approach. This study indicates that smoking, both consciously and unconsciously, contributes to the expansion of asthma and that higher air pollution levels are associated with a higher prevalence of the condition. Furthermore, this study shows that an increased level of connection between smoking and susceptible people leads to a more lasting growth of asthma. We also carry out a numerical analysis to look at the influence of important parameters on the spread of asthma to verify the analytical findings. The main objective of the research is to investigate how smoking affects the prevalence of asthma.

Keywords: Environmental Pollution, Homotopy Perturbation Method (HPM), Asthma Prevalence.

INTRODUCTION

The development of descriptions of systems using mathematical terminology and concepts is known as mathematical modeling. It includes the process of building these models, which are instruments for understanding complex situations, predicting real-world phenomena, and assisting in decision-making. Applications of mathematical modeling can be found in many fields, such as engineering, scientific research, and even the study of





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human behavior. It serves as a link between mathematics and the outside world. Many components of the nonlinear ordinary differential equation system are studied [1]. Fundamental differential equations are receiving a lot of attention these days. They will examine the analytics estimated outcome in this model [2]. Chronic asthma is a respiratory condition marked by inflammation of the airways and an overabundance of mucus. It can cause symptoms like whooping, chest discomfort, coughing, and difficulty inhaling. Numerous things, including allergies, toxins in the environment, and exposure to compounds at work, might cause these symptoms. An ordinary differential equation model was first proposed by Perelson, Kirschner, and De Boer in 1993 [3, 4]. This model is significantly different at first. In many ways, this approach contributed significantly to the non-linear evolution. Stated differently, the nonlinear system beginning and endpoints are included [5,6]. The primary goal is to identify the nonlinear ODE's quantitative solutions [7].By simulating the interactions between these variables over time, mathematical models can help to understand asthma's mechanisms, predict disease progression, and evaluate potential treatment strategies. Such models play a crucial role in advancing our understanding of asthma and guiding clinical interventions. The homotopy perturbation method (HPM) is a powerful mathematical technique used to approximate solutions to a wide range of nonlinear differential equations [8,9]. It involves introducing a small parameter into the equations and expressing the solution as a series expansion in terms of this parameter. Through the systematic perturbation process, the solution is iteratively improved to obtain accurate approximations. An approach for obtaining numerical results is covered [10].

An Approach of Homotopy Perturbation Method

The limits of conventional perturbation methods have been discovered through the application of the HPM approach. Let's examine the non-linear differential equation

$$H_0(k) - f(m) = 0, m \in \phi$$

With boundary conditions

$$B_0(\mathbf{k}, \frac{\partial k}{\partial r}) = 0, r \in \Omega$$
 (1)

when f(m) is an identified mathematical function, B_0 is a boundary function, H_0 is a differential function, as well as ϕ , is a domain with Ω borders. The equation(1) is currently simplified into

$$X(k)+Y(k)-f(m)=0$$

While applying HPM approaches, we obtain

$$D(l,p) = (1-p)[X(l) - X(k_0)] + p[H_0(l) - f(m)] = 0$$

$$D(l,p) = X(l) - X(k_0) + pX(k_0) + P[Y(l) - f(m)] = 0$$
(3)

The origin of equation (1) is k_0 , as well as its embedding parameter $p \in [0,1]$. Applying the limits in the equation (2) and (3), we obtain

$$D(l,0) = X(l) - X(k_0) = 0$$

$$D(l,1) = H_o(l) - f(m) = 0$$

At p=0, the equation (3) gets linear, while at p=1, it turns non-linear. The procedure shall conclude at zero whenever

$$X(l) - X(k_0) = 0$$
 to $H_0(l) - f(m) = 0$

The result is expressed in the following manner: the embedding parameter p remains extremely limited

$$l = l_0 + pl_1 + p^2 l_2 + \dots (4)$$

If p = 1, it represents an approximation to the equation (1)





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$$k = p \xrightarrow{\lim} l = l_0 + l_1 + l_2 + \dots$$
 (5)

In the majority of scenarios, the series converges, Although the non linearfunction H(l) determines the convergent level.

Example

We identify the differential equation of a system that is nonlinear by determining its exact solution

$$z = e^{2y}, z^1 = 2z_2^2 \tag{6}$$

If estimates for both terms in equation (4) are adequate, then we get

$$l = l_0 + pl_1 + p^2 l_2 + \dots (7)$$

If $l_{i,j}$, i.j = 1, 2, 3... are as-yet-undetermined variables. Consequently, using equation (5)

$$z(y) = p \xrightarrow{\lim} l(y) = \sum_{r=0}^{r=2} l_r(y)$$
(8)

The HPM allows us to build a homotopy of equation (6) in the manner described below

$$(1-p)(l-k) + p(l-2l_2^2) = 0 (9)$$

These are the initial estimates
$$l(0) = 1 \& z(0) = 1$$
 (10)

By establishing equations (7)& (10) to equation (9) and remodeling according to the digits of p-terms, the result is

$$\begin{array}{c} \downarrow \\ l_0 + (l_1 - 2l_{2,0}^2) p + (l_2 - 4l_{2,0} 1_{2,1}) p^2 + \dots = 0
\end{array}$$

To derive the unidentified variables $l_{i,j}$, i, j = 1, 2, 3... we need to establish and resolve the system below

$$\vec{l}_0 = 0$$

$$l_2 - 4l_{2.0}l_{2.1} = 0$$

$$l_{0}(v) = 1$$

$$l_1(y) = -e^{-2y} + 8e^{-y} + 8y - 7$$

$$l_2(y) = \frac{1}{3}e^{-4y} - \frac{56}{9}e^{-3y} - 16e^{-2y}y + 6e^{-2y} + \frac{1}{3}e^{-4y} + \frac{272}{3}e^{-y} + 64e^{-y}y + \frac{112}{3}y - \frac{817}{9}e^{-4y}$$

Hence based on equation (8)

$$z(y) = -3 + 6y + 4e^{-y}$$

Mathematical Model

Formulate the system as

$$\frac{dS}{dt} = \wedge -\delta S - \beta SC$$

$$\frac{dI}{dt} = \gamma C - \alpha I - \delta I$$

$$\frac{dC}{dt} = k - \gamma C - \theta C - \delta C$$





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In this model, we divide the population into three categories: Sis the individuals who are susceptible to the disease, Ibe the number of individuals who are infected with a certain disease and C is the individuals who are categorized as smokers. This disease is primarily caused by smoking in public places. By applying the HPM, the system of equations changes to

$$\frac{dS}{dt} - \wedge + \delta S + \beta SC = 0$$

$$\frac{dI}{dt} - \gamma C + \alpha I + \delta I = 0$$

$$\frac{dC}{dt} - k + \gamma C + \theta C + \delta C = 0$$

Using the HPM, the above system can be solved as follows:

$$(1-p)(\frac{dS}{dt} - \lambda + \delta S) + hp(\frac{dS}{dt} + \lambda - \delta S - \beta SC) = 0$$

$$(1-p)(\frac{dI}{dt} - \gamma C + \alpha I + \delta I) + hp(\frac{dI}{dt} + \gamma C - \alpha I - \delta I) = 0$$

$$(1-p)(\frac{dC}{dt} - k + \gamma C + \theta C + \delta C) + hp(\frac{dC}{dt} + k - \gamma C - \theta C - \delta C) = 0$$

Compare the p-coefficient in the system described above

$$\begin{split} P^{0} : & \frac{dS_{0}}{dt} - \wedge + \delta S_{0} = 0 \\ & p^{0} : \frac{dI_{0}}{dt} - \gamma C_{0} + \alpha I_{0} + \delta I_{0} = 0 \\ & p^{0} : \frac{dC}{dt} - k + \gamma C_{0} + \theta C_{0} + \delta C_{0} = 0 \\ & p^{1} : \frac{dS_{1}}{dt} + \delta S_{1} - \frac{dS_{0}}{dt} + \wedge - \delta S_{0} + h \left(\frac{dS_{0}}{dt} + \wedge - \delta S_{0} - \beta S_{0} C_{0} \right) = 0 \\ & p^{1} : \frac{dI_{1}}{dt} - \gamma C_{1} + XI_{1} - \frac{dI_{0}}{dt} + \gamma C_{0} - XI_{0} + h \left(\frac{dI_{0}}{dt} + \gamma C_{0} - XI_{0} \right) = 0 \\ & p^{1} : \frac{dC_{1}}{dt} + YC_{1} - \frac{dC_{0}}{dt} + k - YC_{0} + h \left(\frac{dC_{0}}{dt} + k - YC_{0} \right) = 0 \end{split}$$

The system above requires these initial & boundaries to have an approximate solution

$$S(0) = 0$$
, $I(0) = 0$, $C(0) = 0$ and

$$S(0) = 5500, I(0) = 4000, C(0) = 200$$

Applying initial conditions to the aforementioned system yields

$$S_0 = \frac{\Lambda}{\delta} (1 - e^{-\delta t})$$

$$I_0 = \frac{k\gamma}{XY} (1 - e^{-Xt}) + \frac{k\gamma}{Y(X - Y)} (e^{-Xt} - e^{-Yt})$$





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$$C_0 = \frac{k}{Y} (1 - e^{-Yt})$$

The outcome, when applying the initial conditions, is as follows

$$S_{1} = (5500 + \frac{1}{V^{2}} - \frac{U}{V} \left(\frac{1}{\delta} + \frac{1}{\delta - U} \right)) e^{-\delta t} + \frac{U}{V\delta} + \frac{U e^{-Ut}}{V(\mu - U)} - \frac{Ut e^{-\delta t}}{V} - \frac{e^{-(U + \delta)t}}{V^{2}}$$

$$I_1 = 4000e^{-Xt} + \frac{200\gamma}{(X-Y)}(e^{-Yt} - e^{-Xt})$$

$$C_1 = 200e^{-Yt}$$

Using Homotopy Perturbation Method, we get the approximate solution of the model

$$S(t) = \frac{\wedge}{\delta} (1 - e^{-\delta t}) + (5500 + \frac{1}{V^2} - \frac{U}{V} \left(\frac{1}{\delta} + \frac{1}{\delta - U} \right)) e^{-\delta t} + \frac{U}{V\delta} + \frac{U e^{-Ut}}{V(\mu - U)} - \frac{Ut e^{-\delta t}}{V} - \frac{e^{-(U + \delta)t}}{V^2}$$

$$I(t) = \frac{k\gamma}{XY}(1 - e^{-Xt}) + \frac{k\gamma}{Y(X - Y)}(e^{-Xt} - e^{-Yt}) + 4000e^{-Xt} + \frac{200\gamma}{(X - Y)}(e^{-Yt} - e^{-Xt})$$

$$C(t) = \frac{k}{Y}(1 - e^{-Yt}) + 200e^{-Yt}$$

$$X = \alpha + \delta$$

$$Y = \gamma + \theta + \delta$$

$$U = h\beta \wedge k$$

$$V = (\gamma + \theta + \delta)\delta$$

Numerical Result

We separated people into three sections in order to examine the spread of the asthma illness. We employed numerical results to analyse the impact and spread of asthma in the model. The rate of variation of the susceptible people over duration and various values of β , \wedge and h are examined in figure 1,4 &7.Next, the rate of variation of the diseased people over duration and various values of δ , α and γ are examined in figure 2,5 &8. For figure 3,6 and 9, we examine the behavior of the smoker population over time for various values of δ , γ and θ . Tables 1 and 2 provide a description of the model. More evident, consequently, whether the rate of variation of the diseased people over duration seems high.

CONCLUSION

To analyse how asthma spreads in a population of different type, we used a mathematical model in this study. The model takes into account how smoking affects an individual's risk of developing asthma. An approximate analytical solution for the spread of asthma in a population was found by analysing the model using a mathematical method known as the homotopy perturbation method. It is evident that individuals who are susceptible to infections become infected when exposed to a smoking environment consistently. Numerical models show that when smokers smoke often around others, the rate of variation of harmful pollutants rises over the duration. Therefore, it could be important to restrict the number of smokers entering the population to halt the spread of asthma.





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Table 1: Parameters and Description

Symbol	Description of Parameter
β	Frequency of interaction between smokers and susceptible
δ	Rate of natural death
α	Death rate due to disease
K	The rate at which new individuals become smokers
θ	The frequency of smokers quitting
\wedge	Interaction between pollution and those who are exposed
γ	Rate of infection among smokers

Table 2: Values of the asthma model's parameters

Symbol	The assigned value for a parameter	
^	0.014	
α	0.018	
K	60	
θ	0.002	
$ heta_{\scriptscriptstyle 1}$	0.0002	
β	0.0002	
^	0.0001	





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RESEARCH ARTICLE

A Local and Global Stability Analysis of the Graves Disease Model

G. Lavanya and D. Jasmine

PG & Research Department of Mathematics, Bishop Heber College (Affiliated to Bharathidasan University), Trichy, Tamil Nadu, India.

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*Address for Correspondence

G. Lavanya

PG & Research

Department of Mathematics,

Bishop Heber College (Affiliated to Bharathidasan University),

Trichy, Tamil Nadu, India.

Email: lavanya.maths19@gmail.com



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ABSTRACT

Graves' disease is a thyroid gland autoimmune disorder. We developed a novel mathematical model consisting of four nonlinear ordinary differential equations to describe the disease dynamics. The system of equation consist of variables such as Thyroid Stimulating Hormone (TSH), Free Thyroxine (FT4), thyroid gland volume (V) and TSH receptor autoantibodies (TRAB). The model was constructed by analyzing the underlying biological mechanisms and pathways involved in the pathogenesis of Graves' disease. Key interactions and feedback loops were translated into differential equations based on biochemical reaction kinetics. Analytical stability analysis was then conducted to study the dynamic behavior and equilibrium states of the system. To validate, the model was simulated in simbiology MATLAB. This work establishes a validated mathematical framework capturing the pathogenesis of Graves' disease. By proving and computationally verifying asymptotic stability of the disease state, the model can be further utilized to explore clinical interventions and treatment responses.

Keywords: Thyroid Stimulating Hormone (TSH), Free Thyroxine (FT4), thyroid gland volume (V) and TSH receptor autoantibodies (TRAB).

INTRODUCTION

Graves disease is an autoimmune condition characterized by the hyperactivity of the thyroid gland, causing it to produce excessive amounts of hormones. It ranks among the most prevalent thyroid conditions and stands as the primary cause of hyperthyroidism, a state characterized by the thyroid gland's excessive production of hormones[1,2,3,4]. It is responsible for 60% to 80% of instances of hyperthyroidism. In the US, hyperthyroidism represents 1.2% of the population overall, with an incidence of 20/100,000 to 50/100,000. People between the ages of





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20 and 50 are most likely to have it. According to some research, women are more likely than males to have Graves' disease, with a lifetime risk of 3% compared to 0.5% for men. The 12-year incidence among women between the ages of 25 and 42 was as high as 4.6/1000, according to the Nurses' Health Study II (NHSII)[1]. Hormones that control metabolism are released by the thyroid, a little gland shaped like a butterfly that is located in front of the neck. When the thyroid gland is erratically attacked by the immune system, it generates excessive hormones, which causes a variety of problems throughout the body in Graves disease[2,3]. Once properly diagnosed, Graves' disease is quite manageable. In some instances, it may enter remission or resolve entirely after several months or years. However, if left untreated, it can lead to severe complications, including life-threatening situations. Prompt treatment is crucial to avoid such risks[4,5]. TSH stimulates thyroid follicular cells by attaching to TSH receptors (TSHR), which triggers the thyroid hormones production and dispersion into the bloodstream of triiodothyronine (T3) and thyroxine (T4)[5,6].

The immune system produces antithyroid stimulating receptor antibodies (TRAb) that circulate in the blood and mimic the action of TSH. Consequently, TSH levels drop below the normal range and become almost undetectable in the blood, while TRAb continuously stimulates the gland to produce and secrete hormones, leading to an overactive thyroid gland (hyperthyroidism) in otherwise healthy individuals[7]. Due to a complicated interaction between a genetic trait and environmental factors, Graves' illness is characterised by the development of TRAb and a loss of immunological tolerance to thyroid antigens. Though other organs like the eyes, pituitary, skin, and joints are also included, it is defined as an organ-specific autoimmune disease. Elevated levels of serum free thyroid hormones, specifically free T4 (FT4) and free T3 (FT3), along with undetectable blood TSH and the presence of serum TRAb at some time, are the laboratory confirmation of Graves' hyperthyroidism[7]. A simplified differential equation model were constructed to depict the functioning of the HPT axis in individuals with Hashimoto's thyroiditis. This model comprises four variables: the functional size of the thyroid gland, the concentration of serum TSH, serum FT4, and serum thyroid peroxidase antibody[8]. Similarly, Graves' disease was addressed through a set of differential equations. The initial mechanistic model, designed to predict relapse in Graves' disease patients, was constructed using an ordinary differential equation system to capture the dynamics of the condition[9].

METHODOLOGY

Construction of the Model

The given system of ordinary differential equations (ODEs) represents a mathematical model for Graves' disease, . The model describes the dynamics of thyroid stimulating hormone (TSH), thyroid-stimulating autoantibodies (TRAB), the volume of the thyroid gland (V) and free thyroxine (FT4). The equations governing the rates of change for these variables are as follows:

$$\frac{dTSH}{dt} = k_1 \frac{FT4}{K_a + FT4} - k_2 \frac{TSH}{K_b + TSH} - k_3 TSH + k_4$$

$$\frac{dFT4}{dt} = k_5 V \frac{Ab}{K_c + Ab} - k_6 FT4$$

$$\frac{dV}{dt} = k_7 + k_8 \frac{Ab}{V} - k_9 V$$

$$\frac{dAb}{dt} = k_{prod} - k_{decay} Ab - k_{bind} Ab \frac{TSH}{K_d + TSH}$$
(1)





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Where $k_1, k_2, k_3, k_4, k_5, k_6, k_7, k_8, k_9, k_{prod}, k_{decay}, k_{bind}, K_a, K_b, K_c, K_d > 0$ and TSH(0) > 0, FT4(0) > 0, V(0) > 0 and Ab(0) > 0

In first equation, The term $k_1 \frac{FT4}{K_a + FT4}$ represents the negative feedback regulation of TSH by FT4, where k_1 is

the rate constant, and K_a is the Michaelis-Menten constant for this process. The second term represents the clearance or degradation of TSH, with k_2 as the rate constant and K_b as the Michaelis-Menten constant. The third term accounts for an additional clearance or degradation of TSH, with a rate constant k_3 . The last term k_4 , represents a constant production rate of TSH. In Second equation, the term $k_5 V \frac{Ab}{K_c + Ab}$ represents the production of FT4 stimulated by the thyroid-stimulating autoantibodies (TRAB), where k_5 is the rate constant, V is the volume of the thyroid gland, and K_c is the Michaelis-Menten constant for this process. The second term accounts for the clearance or degradation of FT4, with a rate constant k_6 . In third equation, the first term k_7 , represents a

where k_8 is the rate constant, and the term TRAB/V represents the concentration of TRAB relative to the thyroid gland volume. The third term, $-k_9V$, represents a natural decay or shrinkage of the thyroid gland, with a rate constant k_9 . In fourth equation, the first term, k_{prod} , represents a constant production rate of TRAB. The second term accounts for the natural decay or clearance of TRAB, with a rate constant k_{decay} . The third term represents the binding of TRAB to TSH, which effectively removes TRAB from the system. In this term, k_{bind} is the rate constant, and K_d is the Michaelis-Menten constant for the binding process.

constant widening rate of thyroid gland. The second term accounts for the stimulation of thyroid growth by TRAB,

Theorem 1. Using the initial conditions given in equation (1), the solutions TSH(t), FT4(t), V(t), Ab(t) are non-negative for all time t > 0

Proof:

To prove the non-negativity of TSH(t), let's first analyze the differential equation:

$$\frac{dTSH}{dt} = k_1 \frac{FT4}{K_a + FT4} - k_2 \frac{TSH}{K_b + TSH} - k_3 TSH + k_4$$

Since all the parameters and initial conditions are positive, Now, to prove the non-negativity of TSH(t), By the comparison theorem for differential equations. Let's define a new function $g(t) = e^{k_3 t} TSH(t)$. Taking the derivative of g(t) with respect to t and check its sign:

$$\frac{dg(t)}{dt} = e^{k_3 t} \frac{dTSH}{dt} + k_3 e^{k_3 t} TSH(t)$$

Substituting the given differential equation into the above expression:

$$\frac{dg(t)}{dt} = e^{k_3 t} \left(k_1 \frac{FT4}{K_a + FT4} - k_2 \frac{TSH}{K_b + TSH} - k_3 \cdot TSH + k_4 \right) + k_3 e^{k_3 t} TSH(t)$$

We can simplify this expression:





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$$\frac{dg(t)}{dt} = e^{k_3 t} \left(k_1 \frac{FT4}{K_a + FT4} - k_2 \frac{TSH}{K_b + TSH} + k_4 \right)$$

Now, let's examine the terms in the brackets:

$$1.k_1 \frac{FT4}{Km_1 + FT4} > 0$$
 since all parameters are positive.

2.
$$-k_2 \frac{TSH}{Km_2 + TSH}$$
 because $TSH > 0$ and the other parameters are positive.

3. $k_4 > 0$ as per our assumptions.

Therefore,
$$\frac{dg(t)}{dt} > 0$$
 for all t

Since $\frac{dg(t)}{dt} > 0$, g(t) is an increasing function of t, which implies that TSH(t) is also an increasing function

of t because $e^{k_3 t} > 0$ for all t. Since TSH(0) > 0, and TSH(t) is an increasing function of t, it follows that TSH(t) > 0 for all t. Therefore, we have proved the non-negativity of TSH(t).

Similary, other equations can be proved.

Stability analysis

Theorem 2: The equilibrium E^* is locally asymptotically stable.

Here is an analytical proof of asymptotic stability for the given system of differential equations:

Let's define the state variables

$$x_1 = [TSH], x_2 = [FT4], x_3 = [V], x_4 = [TRAB]$$

Then we can rewrite the system as:

$$\frac{dx_1}{dt} = k_1 \frac{x_2}{K_a + x_2} - k_2 \frac{x_1}{K_b + x_1} - k_3 x_1 + k_4$$

$$\frac{dx_2}{dt} = k_5 x_3 \frac{x_4}{K_c + x_4} - k_6 x_2$$

$$\frac{dx_3}{dt} = k_7 + k_8 \frac{x_4}{x_3} - k_9 x_3$$

$$\frac{dx_4}{dt} = k_{prod} - k_{decay}x_4 - k_{bind}x_4 \frac{x_1}{K_d + x_1}$$

To prove asymptotic stability, we find the equilibrium points by setting dx/dt = 0:





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$$x_{1}^{*} = \frac{\left(k_{1}k_{5}k_{prod} - k_{2}k_{3}K_{a}K_{b}K_{c}\right)}{k_{1}k_{5}k_{6}K_{b}K_{c} + k_{2}k_{3}k_{4}K_{a}K_{c} + k_{1}k_{2}k_{5}K_{a}K_{b}}$$

$$x_{2}^{*} = \frac{k_{5}x_{3}x_{4}^{*}}{k_{6}\left(K_{c} + x_{4}^{*}\right)}$$

$$x_{3}^{*} = \frac{k_{7}}{k_{9}}$$

$$x_{4}^{*} = \frac{k_{prod}}{k_{decay} + \frac{k_{bind}x_{1}^{*}}{K_{d} + x_{1}^{*}}}$$

At the equilibrium point, the Jacobian matrix is evaluated as follows:

$$J = \begin{bmatrix} -\frac{k_2 K_b}{(K_b + x_1^*)^2} - k_3 & \frac{k_1 K_a}{(K_a + x_2^*)^2} & 0 & 0 \\ 0 & -k_6 & \frac{k_5 x_4^*}{K_c + x_4^*} & \frac{k_5 K_c x_3^*}{(K_c + x_4^*)^2} \\ 0 & 0 & -\frac{k_8 x_4^*}{x_3^{*2}} - k_9 & \frac{k_8}{x_3^*} \\ -\frac{K_d k_{bind} x_4^*}{(K_d + x_1^*)^2} & 0 & 0 & -k_{decay} - \frac{k_{bind} x_1^*}{K_d + x_1^*} \end{bmatrix}$$

The trace of the Jacobian matrix is:

$$trace(J) = -k_3 - k_6 - k_9 - k_{decay} - \frac{k_2 K_b}{(K_b + x_1^*)^2} - \frac{k_8 x_4^*}{x_3^{*2}} - \frac{k_{bind} x_1^*}{K_d + x_1^*}$$

Which is negative since all rate constants and concentrations are positive. The determinant of the Jacobian matrix is:

$$\det(J) = \left(\frac{k_2 K_b}{(K_b + x_1^*)^2} + k_3\right) \left(\frac{k_8 x_4^*}{x_3^{*2}} + k_9\right) \left(k_{decay} + \frac{k_{bind} x_1^*}{K_d + x_1^*}\right)$$

$$+ \left(\frac{k_1 K_a}{\left(K_a + x_2^*\right)^2}\right) \left(\frac{K_d k_{bind} x_4^*}{(K_d + x_1^*)^2}\right) \left[\left(\frac{k_5 x_4^*}{K_c + x_4^*}\right) \frac{k_8}{x_3^*} + \left(\frac{k_8 x_4^*}{x_3^{*2}} + k_9\right) \left(\frac{k_5 K_c x_3^*}{\left(K_c + x_4^*\right)^2}\right)\right]$$

$$> 0$$

According to the Routh-Hurwitz stability criterion, the equilibrium point is asymptotically stable since the trace of the matrix is negative and the determinant is positive.





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Therefore, we have analytically proven that the given system of differential equations modeling thyroid hormone regulation is asymptotically stable around its equilibrium point.

Theorem 3. The equilibrium E^* is globally asymptotically stable with some conditions.

Proof. lets denote TSH as T, FT4 as F and Ab as A. let consider the positive definite function

$$U = \frac{1}{2} m_1 (T - T^*)^2 + \frac{1}{2} m_2 (F - F^*)^2 + \frac{1}{2} m_3 (V - V^*)^2 + \frac{1}{2} m_4 (A - A^*)^2$$

Differentiating the above the function with respect to time t, we have

$$\begin{split} \dot{U} &= m_1 \left(T - T^* \right) \dot{T} + m_2 \left(F - F^* \right) \dot{F} + m_3 \left(V - V^* \right) \dot{V} + m_4 \left(A - A^* \right) \dot{A} \\ \dot{U} &= -m_1 \left(\frac{k_2 K_b}{(K_b + T^*)^2} + k_3 \right) \left(T - T^* \right)^2 + m_1 \left(\frac{k_1 K_a}{\left(K_a + F^* \right)^2} \right) \left(T - T^* \right) \left(F - F^* \right) - m_2 (k_6) \left(F - F^* \right)^2 \\ &+ m_2 \left(\frac{k_5 A^*}{K_c + A^*} \right) \left(F - F^* \right) \left(V - V^* \right) + m_2 \left(\frac{k_5 K_c V^*}{\left(K_c + A^* \right)^2} \right) \left(F - F^* \right) \left(A - A^* \right) - \\ &- \left(\frac{k_8 A^*}{V^{*2}} + k_9 \right) m_3 \left(V - V^* \right)^2 + m_3 \frac{k_8}{V^*} \left(V - V^* \right) \left(A - A^* \right) \\ &+ m_4 \left(\frac{K_d k_{bind} A^*}{\left(K_d + T^* \right)^2} \right) \left(A - A^* \right) \left(T - T^* \right) - m_4 \left(k_{decay} + \frac{k_{bind} T^*}{K_d + T^*} \right) \left(A - A^* \right)^2 \end{split}$$

 \dot{U} will be negative definite

$$m_{1} \left(\frac{k_{1}K_{a}}{\left(K_{a}+F^{*}\right)^{2}}\right)^{2} < \frac{2}{3}m_{2}k_{6}k_{3}$$

$$m_{2} \left(\frac{k_{5}A^{*}}{K_{c}+A^{*}}\right)^{2} < \frac{2}{3}m_{3}k_{6}k_{9}$$

$$m_{4} \left(\frac{K_{d}k_{bind}A^{*}}{\left(K_{d}+T^{*}\right)^{2}}\right)^{2} < \frac{2}{3}m_{1}k_{3}k_{decay}$$

$$m_{2} \left(\frac{k_{5}K_{c}V^{*}}{\left(K_{c}+A^{*}\right)^{2}}\right)^{2} < \frac{4}{9}m_{4}k_{6}k_{decay}$$

$$m_{3} \left(\frac{k_{8}}{V^{*}}\right)^{2} < \frac{2}{3}m_{4}k_{9}k_{decay}$$





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Now $m_1 = 1$, $m_3 = 1$, $m_4 = 1$

$$\left(\frac{k_1 K_a}{\left(K_a + F^*\right)^2}\right)^2 < \frac{4}{9} k_6 \left(K_c + A^*\right)^2 \min\left\{\frac{k_9}{k_5 (A^*)^2}, \frac{2}{3} \frac{k_{decay}}{K_c V^*}\right\}$$

 \dot{U} will be negative definite provided the above conditions are satisfied and hence E^* is globally asymptotically stable.

RESULTS AND DISCUSSION

Graves' disease involves a characteristic progression from normal thyroid hormone levels (euthyroidism) to overt hyperthyroidism. This progression typically follows a pattern marked by a decline in TSH (thyroid stimulating hormone) levels and an increase in FT4 (free thyroxine) levels. The drop in TSH is caused by negative feedback from the elevated circulating levels of thyroid hormones FT4 and FT3 (free triiodothyronine)[10,11]. Meanwhile, the rise in FT4 results from the overproduction of thyroid hormones by the overactive thyroid gland in Graves' disease. So the TSH decreases in response to the excess thyroid hormones, while the FT4 increases due to the gland's excessive hormone production and release. Let the Initial values of TSH 0.906 mU/L[13], FT4 36 pg/mL[7], thyroid volume (V) 30 mL[7,9], and antibody levels (Ab) 25 IU/mL[7,9] for the parameters values from the table.

We can plug in the initial conditions into the TSH rate equation:

$$0 = k_1(36)/(0.0434+36) - k_2(0.906)/(K_a+0.906) - 16.6355(0.906) + 30$$

$$k_1 = 15.0479 + k_2(0.906)/(K_a+0.906)$$

Lets assume:

$$k_2 = 0.5 / day$$
, $K_a = 1 mU/L$ (guess)

$$k_1 = 15.0479 + 0.5(0.906)/(1+0.906) = 15.45 \, mU/L \, per \, (pg/mL \, x \, day)$$

In some cases, certain parameter values are calculated using the initial condition and known parameter values. For the remaining parameter values, an optimization approach is employed using the fmincon function in MATLAB. The upper and lower bounds for these parameters are set based on values reported in previous literature studies[10,12]. This optimization process simulates or estimates the unknown parameter values by finding the optimal solution within the specified bounds, leveraging the known information and the initial conditions.

CONCLUSION

A novel mathematical framework has been established to elucidate the pathophysiology of Graves' disease. This framework, formulated as a system of four nonlinear ordinary differential equations, integrates critical disease variables including thyroid-stimulating hormone (TSH), free thyroxine (FT4), thyroid gland volume (Fsize), and TSH receptor autoantibodies (TRAB). The development meticulously incorporates established biological mechanisms underlying Graves' disease pathogenesis. The system's dynamic behavior and equilibrium states were investigated through analytical stability analysis. Additionally, Lyapunov theory was employed to mathematically demonstrate the model's asymptotic stability. Parameter estimation utilized a combined approach, leveraging both clinical data and optimization techniques implemented using fmincon within the MATLAB environment. To ensure validity, the model underwent comprehensive simulation within simbiology MATLAB. This work represents a significant





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contribution by establishing a validated mathematical framework that captures the essential features of Graves' disease pathogenesis. The model's utility extends beyond mere comprehension, it serves as a powerful tool to explore potential clinical interventions and analyze treatment responses. In conclusion, this research offers a valuable contribution to the field of Graves' disease by providing a robust mathematical modeling approach, paving the way for future investigations into optimized treatment strategies.

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Table 1: Normal range and disease state

	Normal Range	Disease state	Source	Unit
TSH	0.4 - 4.0	< 0.4	Link(verywellhealth)	mU/L
FT4	7-18	>18	Literature(Mo C et al, 2023)	Pg/mL
V	10-18	-	Literature(Sakane 1990)	mL
Ab	0-1.75	>1.75	Literature(Mo C et al, 2023)	IU/L

Table 2: parameters description and values with units

	description	values	Units
k_1	Rate constant for TSH stimulation by FT4	15.45	mU/L*day
k_2	Rate constant for TSH auto-inhibition	0.5	mU/L*day
k_3	Rate constant for TSH degradation	16.6355	$pg / mL^2 * day$





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k_4	Basal TSH production rate	30	$pg / mL^2 * day$
k_5	Rate of FT4 production stimulated by TSH and functional size of thyroid gland		-
k_6	Rate constant for FT4 degradation	0.0183	$ml^3/U*day$
k_7	constant widen rate of thyroid gland.	0.0289	-
k_8	Constant widen rate of thyroid gland stimulated by TRAb	0.2781	U / mL*day
k_9	constant decay or shrinkage rate of the thyroid gland	0.0316	U / mL*day
k_{prod}	Production rate constant for TRAb	3.1052	1 / day
k_{decay}	Decay/clearance rate of TRAb	0.0087	-
k_{bind}	Michaelis-Menten constant for TRAb binding to TSH.	0.0866	mU/L
K_a	Michaelis constant for FT4 stimulation of TSH	0.0434	1 / <i>day</i>
K_b	Michaelis constant for TSH auto-inhibition	1	1 / day
K_c	Michaelis constant for TSH stimulation of FT4	0.004	1 / day
K_d	Michaelis constant for TRAB binding to TSH	1	1 / day

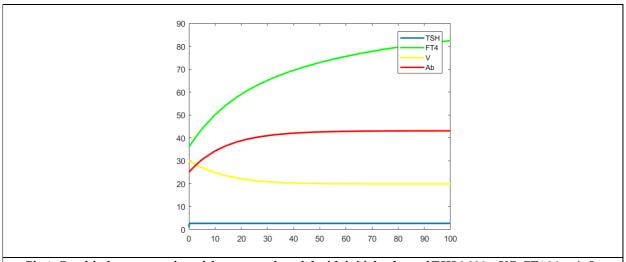


Fig 1: Graphical representation of the proposed model with initial values of TSH 0.906 mU/L, FT4 36 pg/mL, thyroid volume (V) 30 mL, and antibody levels (Ab) 25 IU/mL for the parameters values from the table 2.





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RESEARCH ARTICLE

Some New Results on Harary Energy of Graphs

M. Deva saroja¹ and V. Vinisha²

¹Assistant Professor, Rani Anna Government College for Women, (Affliated to Manonmaniam Sundaranar University), Tamil Nadu, India.

²Research Scholar, Registration Number: 23111172092002, Rani Anna Government College for Women, (Affliated to Manonmaniam Sundaranar University) Tamil Nadu, India.

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*Address for Correspondence

M. Deva saroja

Assistant Professor,

Rani Anna Government College for Women,

(Affliated to Manonmaniam Sundaranar University),

Tamil Nadu, India.

Email: mdsaroja@gmail.com



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ABSTRACT

In this paper, we compute the Harary energy of some graphs. Also, we obtained the Harary energy of regular graph obtained from join of complete graphs.

Keywords: Harary matrix, Harary energy, cocktail party graph, crown graph, friendship graph. 2020 Mathematics Subject Classification: 05C50

INTRODUCTION

Throughout the paper, all graphs wil be assumed to be simple and finite in order to simplify the analysis. There has been a lot of discussion about the concept of Energy of graph since 1978, when Ivan Gutman introduced it. A more detailed explanation of the concept energy can be found in [2,3]. The focus of this article is on Harary matrix[4], developed by Ivanciuc et al [5] in the 1970s. Assume that G is an undirected graph consisting of a vertices set $V(G) = \{n_1, n_2, ..., n_p\}$ and edges set E(G). In the case of this vertices n_i and n_k , the distance between them, denoted by D'_{jk} , can be considered to the length of the shortest path connecting this two vertices. The Harary matrix[4] $H(G) = [R_{jk}]$ of G is a $p \times p$ matrix, where

$$R_{jk} = \begin{cases} \frac{1}{D'_{jk}}, & if \ j \neq k \\ 0, & if \ j = k \end{cases}$$





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A group of eigen values from H(G) denoted by $\eta_1 \ge \eta_2 \ge \cdots \ge \eta_p$ are known as Harary eigenvalues and their collection is known as Harary spectrum of G. For more details on Harary eigenvalues, refer [6,7,8]. Harary energy of a graph G, referred to as $E_H(G)$, can be defined [9] as follows:

$$E_H(G) = \sum_{i=1}^p |\eta_i|$$

H(G) is defined to have the characteristic polynomial of the form $P(G,\eta) = \det(\eta I - H(G))$. In this, I represents the unit matrix of order p. The eigen values of H(G) are the roots of $P(G, \eta)$. Since, the Harary matrix H(G) be a real symmetric with zero trace, the collection η_i 's must be real with $\Sigma \eta_i = 0$.

PRELIMINARIES

Lemma 2.1. [10] If U, V, X and Y are all matrices with U non-singular, then we can write $\begin{vmatrix} U & V \\ V & V \end{vmatrix} = |U||Y - XU^{-1}V|$ **Lemma 2.2.** [10] If U, V, X and Y are all matrices. Let $R = \begin{pmatrix} U & V \\ X & V \end{pmatrix}$ if U and X are commutative matrices, then |R| = |UY - XV|

Lemma 2.3. [11] The adjacency matrix $A(K_q)$ of complete graph K_q , then $A^2(K_q) = (q-2)A(K_q) + (q-1)I_q$.

Definition 2.4. [12] Let us take a pair of complete graphs K_q with vertices set $\{p_i, i=1,2,3,...,q\}$ and $\{r_j, j=1,2,3,...,q\}$ 1,2,3,...,q . Construct a graph by joining p_i to r_i , for i=1,2,3,...q and denote it by $J(K_q^q)$. It has 2q vertices and q^2 edges and is *q*-regular.

HARARY ENERGY OF SOME GRAPHS

Theorem 3.1. Let $\mathit{CP'}_{2q}$ be the cocktail party graph. Then

$$S_pH(CP'_{2q}) = \begin{bmatrix} \left(\frac{4q-3}{2}\right) & \left(\frac{-3}{2}\right) & \left(\frac{-1}{2}\right) \\ 1 & q-1 & q \end{bmatrix}$$

and the Harary energy of CP'_{2q} is, $E_H(CP'_{2p}) = 4q - 3$ where $q \ge 2$

Proof: Let the cocktail party graph be CP'_{2q} then it has 2q vertices, 2q(q-1) edges and is (2q-2) - regular. Let $\alpha = \{\alpha_1, \alpha_2, \alpha_3, \dots, \alpha_{2q}\} \text{ be the eigen values of Harary matrix of } CP'_{2q}. \text{ Then the Harary matrix of } CP'_{2q} \text{ has the form } H(CP'_{2q}) = \begin{bmatrix} A(K_q) & \frac{1}{2}I_q + A(K_q) \\ \frac{1}{2}I_q + A(K_q) & A(K_q) \end{bmatrix}$

$$H(CP'_{2q}) = \begin{bmatrix} A(K_q) & \frac{1}{2}I_q + A(K_q) \\ \frac{1}{2}I_q + A(K_q) & A(K_q) \end{bmatrix}$$

The corresponding characteristic polynomial is

e corresponding characteristic polynomial is
$$|\alpha I_{2q} - H(CP'_{2q})| = \begin{vmatrix} \alpha I_q - A(K_q) & -\left(\frac{1}{2}I_q + A(K_q)\right) \\ -\left(\frac{1}{2}I_q + A(K_q)\right) & \alpha I_q - A(K_q) \end{vmatrix}$$

$$= \left| \left(\alpha I_q - A(K_q)\right)^2 - \left(\frac{1}{2}I_q + A(K_q)\right)^2 \right| \text{ by lemma (2.2)}$$

$$= \left| \left(\alpha^2 - \frac{1}{4}\right)I_q - (2\alpha + 1)A(K_q) \right|$$

$$= (2\alpha + 1)^q \left| \frac{(\alpha^2 - \frac{1}{4})}{(2\alpha + 1)}I_q - A(K_q) \right|$$

$$= (2\alpha + 1)^q \left(\frac{2\alpha - 1}{4} - (q - 1)\right) \left(\frac{2\alpha - 1}{4} + 1\right)^{q - 1}$$

$$= \left(\alpha - \left(\frac{-1}{2}\right)\right)^q \left(\alpha - \left(\frac{4q - 3}{2}\right)\right) \left(\alpha - \left(\frac{-3}{2}\right)\right)^{q - 1}$$

Hence, the Harary spectrum of CP'_{2q} is





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$$S_pH(D_1(K_{2q})) = \begin{bmatrix} \left(\frac{4q-3}{2}\right) & \left(\frac{-3}{2}\right) & \left(\frac{-1}{2}\right) \\ 1 & q-1 & q \end{bmatrix}$$
 and the Harary energy of CP'_{2p} is, $E_H(CP'_{2q}) = 4q-3$ where $q \geq 2$

Theorem 3.2. Let $H'_{q,q}$ be the crown graph. Then

$$S_{p}H(H'_{q,q}) = \begin{bmatrix} \left(\frac{-3q+1}{6}\right) & \frac{9q-7}{6} & \frac{1}{6} & \left(\frac{-7}{6}\right) \\ 1 & 1 & q-1 & q-1 \end{bmatrix}$$

and the Harary energy is, $E_H(H'_{q,q}) = \frac{2(5q-4)}{2}$ where $q \ge 3$

Proof: Let $H'_{q,q}$ be the crown graph then it has 2q vertices, q(q-1) edges and is (q-1) –regular. Let $\eta =$

Proof: Let
$$H_{q,q}$$
 be the crown graph then it has $2q$ vertices, $q(q-1)$ edge $\{\eta_1,\eta_2,\eta_3,...,\eta_{2q}\}$ be the eigen value. Then, the Harary matrix of $H'_{q,q}$ is
$$H(H'_{q,q}) = \begin{bmatrix} \frac{1}{2}A(K_q) & \frac{1}{3}I_q + A(K_q) \\ \frac{1}{3}I_q + A(K_q) & \frac{1}{2}A(K_q) \end{bmatrix}$$

By lemma (2.3), we get the Characteristics polynomia

$$\left(\eta - \left(\frac{-3q+1}{6}\right)\right) \left(\eta - \left(\frac{9q-7}{6}\right)\right) \left(\eta - \frac{1}{6}\right)^{q-1} \left(\eta - \left(\frac{-7}{6}\right)\right)^{q-1}$$
to the spectrum

$$S_{p}H(H_{q,q}) = \begin{bmatrix} \left(\frac{-3q+1}{6}\right) & \frac{9q-7}{6} & \frac{1}{6} & \left(\frac{-7}{6}\right) \\ 1 & 1 & q-1 & q-1 \end{bmatrix}$$

and the Harary energy is, $E_H(H'_{q,q}) = \frac{2(5q-4)}{3}$ where $q \ge 3$

Theorem 3.3. The complete bipartite graph $K_{q,q}$ has the spectru

$$S_p H(K_{q,q}) = \begin{bmatrix} \left(\frac{-1}{2}\right) & \frac{-(q+1)}{2} & \frac{3q-1}{2} \\ 2(q-1) & 1 & 1 \end{bmatrix}$$

and the Harary energy of $K_{q,q}$ is, $E_H(K_{q,q}) = 3q - 1$ when

Proof: Let the complete bipartite graph be $K_{q,q}$ and the eigen value of Harary matrix are $\gamma = \{\gamma_1, \gamma_2, \gamma_3, ..., \gamma_{2q}\}$ respectively. Then, the Harary matrix of $K_{q,q}$ is

$$H(K_{q,q}) = \begin{bmatrix} \frac{1}{2}A(K_q) & I_p + A(K_q) \\ I_q + A(K_q) & \frac{1}{2}A(K_q) \end{bmatrix}$$

By lemma (2.3), we get

$$\left|\gamma I_{2q} - H(K_{q,q})\right| = \left(\gamma + \frac{1}{2}\right)^{2(q-1)} \left(\gamma + \left(\frac{q+1}{2}\right)\right) \left(\gamma - \left(\frac{3q-1}{2}\right)\right)$$

$$S_pH(K_{q,q}) = \begin{bmatrix} \left(\frac{-1}{2}\right) & \frac{-(q+1)}{2} & \frac{3q-1}{2} \\ 2(q-1) & 1 & 1 \end{bmatrix}$$
 and $E_H(K_{q,q}) = 3q-1$ where $q \ge 2$

Theorem 3.4. Let the complete graph be K_a . Then

$$S_p H(K_q) = \begin{bmatrix} (-1) & q-1 \\ q-1 & 1 \end{bmatrix}$$

 $S_pH(K_q) = \begin{bmatrix} (-1) & q-1 \\ q-1 & 1 \end{bmatrix}$ and the Harary energy of K_q is, $E_H(K_q) = 2(q-1)$ where $q \ge 2$

Proof: Let $\beta' = \{\beta'_1, \beta'_2, \beta'_3, ..., \beta'_q\}$ be the eigen value of complete graph K_q . Then it has q vertices, $\frac{q(q-1)}{2}$ edges and is (q-1)- regular. Then the Harary matrix of K_q has the form $H(K_q) = [A(K_q)].$





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By lemma (2.3), we get

$$(\beta'+1)^{q-1}\bigl(\beta'-(q-1)\bigr)$$

Hence,

$$S_p H(K_q) = \begin{bmatrix} (-1) & q-1 \\ q-1 & 1 \end{bmatrix}$$

 $S_pH(K_q)=\begin{bmatrix} (-1) & q-1\\ q-1 & 1 \end{bmatrix}$ and the Harary energy of K_q is, $E_H(K_q)=2(q-1)$ where $q\geq 2$

Theorem 3.5. For any $q \ge 1$, the star graph $K_{1,q}$ has the spectrum

$$S_{p}H(K_{1,q}) = \begin{bmatrix} \left(\frac{-1}{2}\right) & \frac{\left(\frac{q-1}{2}\right) + \sqrt{\left(\frac{q-1}{2}\right)^{2} + 4q}}{2} & \frac{\left(\frac{q-1}{2}\right) - \sqrt{\left(\frac{q-1}{2}\right)^{2} + 4q}}{2} \\ q - 1 & 1 & 1 \end{bmatrix}$$

and the Harary energy of $K_{1,q}$ is, $E_H(K_{1,q}) = \left(\frac{q-1}{2}\right) + \sqrt{\left(\frac{q-1}{2}\right)^2 + 4q}$

Proof: Let us take a star graph $K_{1,q}$ with vertices set $V' = \{v'_0, v'_1, v'_2, ..., v'_q\}$ where the vertex v'_0 has degree q and the eigen values of Harary matrix are $\mu = \{\mu_0, \mu_1, ..., \mu_q\}$ respectively. For this graph, the Harary matrix is $\begin{bmatrix} 0 & 1 & 1 & 1 & \cdots & 1 & 1 \\ 1 & 0 & 1 & 0 & 1 & 1 \end{bmatrix}$

$$H(K_{1,q}) = \begin{bmatrix} 0 & 1 & 1 & 1 & 1 & \cdots & 1 & 1 \\ 1 & 0 & 1/2 & 1/2 & \cdots & 1/2 & 1/2 \\ 1 & 1/2 & 0 & 1/2 & \cdots & 1/2 & 1/2 \\ 1 & 1/2 & 1/2 & 0 & \cdots & 1/2 & 1/2 \\ \vdots & \vdots & \vdots & \vdots & \ddots & \vdots & \vdots \\ 1 & 1/2 & 1/2 & 1/2 & \cdots & 0 & 1/2 \\ 1 & 1/2 & 1/2 & 1/2 & \cdots & 1/2 & 0 \end{bmatrix}$$

On simplification, we get the characteristic polynomial

$$\left(\mu + \frac{1}{2}\right)^{q-1} \left(\mu - \left(\frac{\left(\frac{q-1}{2}\right) \pm \sqrt{\left(\frac{q-1}{2}\right)^2 + 4q}}{2}\right)\right)$$
 and the spectrum would be

$$S_pH(K_{1,q}) = \begin{bmatrix} \left(\frac{-1}{2}\right) & \frac{\left(\frac{q-1}{2}\right) + \sqrt{\left(\frac{q-1}{2}\right)^2 + 4q}}{2} & \frac{\left(\frac{q-1}{2}\right) - \sqrt{\left(\frac{q-1}{2}\right)^2 + 4q}}{2} \\ q - 1 & 1 & 1 \end{bmatrix}$$

Therefore,

$$E_H(K_{1,q}) = \left(\frac{q-1}{2}\right) + \sqrt{\left(\frac{q-1}{2}\right)^2 + 4q}$$

Theorem 3.6. The spectrum and energy of the friendship graph
$$F_q^3$$
 is
$$S_pH(F_q^3) = \begin{bmatrix} 0 & -1 & \frac{q+\sqrt{q^2+8q}}{2} & \frac{q-\sqrt{q^2+8q}}{2} \\ q-1 & q & 1 & 1 \end{bmatrix}$$

and $E_H(F_q^3) = q + \sqrt{q^2 + 8q}$

Proof: Consider the friendship graph F_q^3 with vertices set $V' = \{v'_0, v'_1, v'_2, ..., v'_{2q}\}$ where $\deg(v'_0) = 2q$ and the eigen values of Harary matrix are $\eta' = \{\eta'_0, \eta'_1, ..., \eta'_{2q}\}$ respectively. Then the Harary matrix of F_q^3 is $\begin{bmatrix} 0 & 1 & 1 & 1 & 1 & 1 & 1 \\ 1 & 0 & 1 & 1 & 1 & 1 \end{bmatrix}$

$$H(F_q^3) = \begin{bmatrix} 0 & 1 & 1 & 1 & 1 & \cdots & 1 & 1 \\ 1 & 0 & 1 & 1/2 & 1/2 & \cdots & 1/2 & 1/2 \\ 1 & 1 & 0 & 1/2 & 1/2 & \cdots & 1/2 & 1/2 \\ 1 & 1/2 & 1/2 & 0 & 1 & \cdots & 1/2 & 1/2 \\ 1 & 1/2 & 1/2 & 1 & 0 & \cdots & 1/2 & 1/2 \\ \vdots & \vdots & \vdots & \vdots & \vdots & \ddots & \vdots & \vdots \\ 1 & 1/2 & 1/2 & 1/2 & 1/2 & \cdots & 0 & 1 \\ 1 & 1/2 & 1/2 & 1/2 & 1/2 & \cdots & 1 & 0 \end{bmatrix}$$





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The characteristic polynomial becomes

$$\eta^{q-1}(\eta+1)^q \left(\eta - \left(\frac{q \pm \sqrt{q^2 + 8q}}{2}\right)\right)$$

Hence, the spectrum

$$S_p H(F_q^3) = \begin{bmatrix} 0 & -1 & \frac{q + \sqrt{q^2 + 8q}}{2} & \frac{q - \sqrt{q^2 + 8q}}{2} \\ q - 1 & q & 1 & 1 \end{bmatrix} \text{ and } E_H(F_q^3) = q + \sqrt{q^2 + 8q}$$

Harary energy of regular graph obtained from join of complete graph

Theorem 4.1. Let $J(K_q^q)$ be the join of complete graph. Then the Harary spectrum of $J(K_q^q)$ is

$$S_p H(J(K_q^q)) = \begin{bmatrix} \left(\frac{3q-1}{2}\right) & \frac{q-3}{2} & \left(\frac{-1}{2}\right) & \left(\frac{-3}{2}\right) \\ 1 & 1 & q-1 & q-1 \end{bmatrix}$$

and the Harary energy of $J(K_q^q)$ is $E_H(J(K_q^q)) = 4(q-1)$; $q \ge 3$

Proof: Let $J(K_q^q)$ be the join of complete graph of order 2q, q = 3,4,...,n and q^2 edges. Let the eigen value of Harary matrix be $= \{\lambda_1, \lambda_2, \lambda_3, ..., \lambda_{2q}\}$. Then the Harary matrix of $J(K_q^q)$ has the form

$$H\left(J\left(K_q^q\right)\right) = \begin{bmatrix} A(K_q) & I_q + \frac{1}{2}A(K_q) \\ I_q + \frac{1}{2}A(K_q) & A(K_q) \end{bmatrix}$$

By lemma (2.3), we get the characteristic polynomia

$$\begin{aligned} \left| \lambda I_{2q} - H \left(J(K_q^q) \right) \right| &= \begin{vmatrix} \lambda I_q - A(K_q) & -\left(I_q + \frac{1}{2}A(K_q) \right) \\ -\left(I_q + \frac{1}{2}A(K_q) \right) & \lambda I_q - A(K_q) \end{vmatrix} \\ &= \left| (\lambda I_q - A(K_q))^2 - (I_q + \frac{1}{2}A(K_q))^2 \right| \text{ by lemma (2.2)} \\ &= \left| (\lambda^2 - 1)I_q + \frac{3}{4}A^2(K_q) - (2\lambda + 1)A(K_q) \right| \\ &= \left| (\lambda^2 - 1)I_q + \frac{3}{4}((q - 2)A(K_q) + (q - 1)I_q) - (2\lambda + 1)A(K_q) \right| \text{ by lemma (2.3)} \\ &= \left(\lambda^2 + \frac{3q}{4} - \frac{7}{4} \right) I_q - \left(2\lambda - \frac{3q}{4} + \frac{10}{4} \right) A(K_q) \end{aligned}$$

Simplifying, we get

$$\left(\lambda - \left(\frac{3q-1}{2}\right)\right) \left(\lambda - \left(\frac{q-3}{2}\right)\right) \left(\lambda + \frac{1}{2}\right)^{q-1} \left(\lambda + \frac{3}{2}\right)^{q-1}$$

Hence, the Harary spectrum of $J(K_a^q)$ is

$$S_pH(J(K_q^q)) = \begin{bmatrix} \left(\frac{3q-1}{2}\right) & \frac{q-3}{2} & \left(\frac{-1}{2}\right) & \left(\frac{-3}{2}\right) \\ 1 & 1 & q-1 & q-1 \end{bmatrix} \text{ and } E_H(J(K_q^q)) = 4(q-1) \text{ where } q \ge 3$$

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RESEARCH ARTICLE

Parameter Uniform Convergence of a Finite Element Method for a Singularly Perturbed Linear Reaction Diffusion System Discontinuous Source Terms and Same Perturbation Parameter

Vinoth Maruthamuthu and Joseph Paramasivam Mathiyazhagan*

Department of Mathematics, Bishop Heber College, (Affiliated to Bharathidasan University), Tiruchirappalli, Tamil Nadu, India.

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*Address for Correspondence Joseph Paramasivam Mathiyazhagan

Department of Mathematics, Bishop Heber College, (Affiliated to Bharathidasan University), Tiruchirappalli, Tamil Nadu, India. Email: paramasivam.ma@bhc.edu.in



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ABSTRACT

A linear system of 'n' second order ordinary differential equations of reaction diffusion type with discontinuous source terms and same perturbation parameter is considered. On a piecewise uniform Shishkin mesh, a numerical system is built that employs the finite element method. The numerical approximations obtained by this approach are proven to be effectively almost second order convergent.

Keywords: singular perturbation problems, system of differential equations, reaction diffusion equations, overlapping boundary and interior layers, finite element method, Shishkinmesh, parameter uniform convergence, discontinuous source terms.

INTRODUCTION

Singularly perturbed differential equations are found in many disciplines of applicable mathematics. Many scholars have focused on the analytical and numerical treatment of these equations [1-22]. In general, simple numerical approaches do not give satisfactory approximations for these equations. As a result, one must seek for non-classical ways. Over the last three decades, there have been several publications published on non-classical approaches, the majority of which deal with second order problems. However, only a few authors have produced numerical approaches for singularly perturbed systems of ordinary differential equations [6-22]. In contrast to the work cited in reference [13], all singularly perturbed parameters remain consistent throughout our present paper. This consistency ensures a coherent framework for analysis and comparison across different sections or examples within our study.





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By maintaining uniformity in these parameters, we aim to facilitate clearer and more accurate interpretations of our results and methodologies. In our current paper, inspired by the work of Miller [6], we delve into the discussion of approximate solutions generated through numerical methods. We emphasize the necessity for these solutions to be globally established at every point across the domain of the exact solution, particularly when representing a boundary layer with the chosen numerical approach. A fundamental consideration lies in the utilization of interpolation techniques to achieve this global establishment. Specifically, we advocate for the adoption of basic interpolation methods, such as piecewise linear interpolation. This approach facilitates the extension of the numerical solution beyond mesh points, thereby ensuring representation throughout the entire domain. Given our aspirations to tackle complex scenarios in higher dimensions, we opt to focus on finite-element subspaces utilizing piecewise polynomial basis functions. This choice enables us to maintain flexibility and precision in approximating solutions across diverse and intricate domains. A linear system of 'n' second order ordinary differential equations of the reaction diffusion type with discontinuous source terms is examined in the interval as a singularly perturbed system. Assume that there is just one discontinuity in the source terms at the position $d \in \Omega$. The jump at d in any function $\vec{\phi}$ is defined by $|\vec{\phi}|(d) = \vec{\phi}(d+) - \vec{\phi}(d-)$. We look at the equation that follows

$$-E\vec{y}''(x) + A(x)\vec{y}(x) = \vec{f}(x) \ on\Omega^- \cup \Omega^+, \tag{1}$$

 \vec{y} given on{0,1}and $\vec{f}(d+) \neq \vec{f}(d-)$, where $\Omega^- = \{x: 0 < x < d\}$, $\Omega^+ = \{x: d < x < 1\}$. Here \vec{y} is a column n vector, E and E and E and E are assumed to satisfy components E are assumed to satisfy components E and E are assumed to satisfy E are assumed to satisfy E are assumed to satisfy E and E are assumed to satisfy E are assumed to satisfy E and E are assumed to satisfy E are assumed to satisfy E and E are assumed to satisfy E and E are assumed to satisfy E are assumed to satisfy E and E are assumed to satisfy E are assumed to satisfy E and E are assumed to satisfy E and E are assumed to satisfy E are assumed to satisfy E and E are assumed to satisfy E are assumed to satisfy E and E are assumed to satisfy E and E are assumed to satisfy E and E are assumed to satisfy E and E are assumed to satisfy E and E are assumed to satisfy E and E are assumed to satisfy E and E

$$a_{ii}(x) > \sum_{\substack{j \neq i \\ j=1}}^{n} |a_{ij}(x) + b_i(x)| \text{ for } 1 \le i \le n \text{ and } a_{ij}(x), b_i(x) \le 0 \text{ for } i \ne j$$
 (2)

and, for some α ,

$$0 < \alpha < \min_{\substack{x \in [0,2] \\ 1 \le i \le n}} \sum_{j=1}^{n} |a_{ij}(x) + b_i(x)|. \tag{3}$$

EVALUATION OF THE FINITE ELEMENT MODEL

Let V a given Hilbert space with scalar product (\cdot,\cdot) and a norm of $\|\cdot\|_V$. V is usually a subspace of the Sobolev space $H_0^1(\Omega)$. Examine the weak formulation, find $\vec{y} \in H_0^1(\Omega)^n$ in particular $y_i \in H_0^1(\Omega^- \cup \Omega^+)$ for i = 1, ..., n such that

$$\beta_i(y_i(x), z_i(x)) = f_i(v_i)(x) \,\forall \, z_i(x) \in H_0^1(\Omega^- \cup \Omega^+)$$

$$f_i(v_i)(x) = (f_i(x), v_i(x)),$$
(4)

where $(y_i(x), z_i(x)) = \int y_i(x)z_i(x)dx$.

$$\beta_i(y_i(x), z_i(x)) = -\varepsilon \quad (y_i(x), z_i(x)) + \left(\sum_{j=1}^n (a_{ij}(x)y_j(x)), z_i(x)\right),$$

 $f_i(z_i(d+)) \neq f_i(z_i(d-)), f_i(z_i)(x)$ is a continuous linear functional on $H_0^1(\Omega^- \cup \Omega^+)^n$ and $\beta_i(y_i(x), z_i(x))$ are bilinear forms on $H_0^1(\Omega^- \cup \Omega^+)^n$.

Lemma 1 Suppose that the bilinear forms $\beta_i(y_i(x), z_i(x))$, i = 1, ..., n, is continuous on $H_0^1(\Omega^- \cup \Omega^+)^n$ is coercive, that

$$\left|\beta_{i}(y_{i}(x), z_{i}(x))\right| \leq \gamma_{1} \|y_{i}(x)\| \|z_{i}(x)\| \tag{5}$$

 $\beta_i(z_i(x), z_i(x)) \ge \alpha \parallel z_i(x) \parallel^2 \tag{6}$

The constants α and γ_1 are independent of y_i and z_i . For each continuous linear functional $f_i(\cdot)$, problem (4) has a unique solution.

Lemma 2 Satisfies the coercive property with respect to the bilinear functional $\beta_i(y_i(x), z_i(x))$, i = 1, ..., n, $\|z_i\|_{\mathcal{E}}^2 \leq \beta_i(z_i, z_i)$





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Proof: Each i, i = 1, ..., n

$$\beta_{i}(z_{i}, z_{i}) = -\varepsilon \left(z_{i}^{'}, z_{i}^{'}\right) + \left(\sum_{j=1}^{n} (a_{ij}z_{j}), z_{i}\right)$$

$$= \varepsilon \quad \|z_{i}\|_{1}^{2} + \int_{0}^{1} \left(\sum_{j=1}^{n} (a_{ij}z_{j}) \cdot z_{i}\right) dx$$

$$\geq \varepsilon \quad \|z_{i}\|_{1}^{2} + \alpha \|z_{i}\|_{0}^{2}.$$

FORMATION OF THE SHISHKIN MESH

We now build a piecewise uniform Shishkin mesh on $\Omega^- \cup \Omega^+$ with N mesh-intervals. Let $\Omega^N = \Omega^{-N} \cup \Omega^{+N}$ where $\Omega^{-N} = \{x_k\}_{k=1}^{\frac{N}{2}-1}$, $\Omega^{+N} = \{x_k\}_{k=0}^{N-1}$. The mesh $\overline{\Omega}^N$, which is piecewise uniform on [0,1], is produced by splitting [0,d] into three mesh-intervals, as shown below:

produced by splitting [0,d] into three mesh-intervals, as shown below:
$$\sigma = \min \left\{ \frac{d}{4}, \frac{2\sqrt{\varepsilon}}{\sqrt{\alpha}} \ln N \right\}$$
[0,\sigma] \(\cup (\sigma , d - \sigma) \cup (d - \sigma , d)\)
(7)

Next, on the sub-interval(σ , $d-\sigma$], a uniform mesh of N/4 mesh-points is created; on each of the sub-intervals $(0,\sigma]$ and $(d-\sigma)$, d a uniform mesh of N/8 mesh-points is set. The remainder is produced by splitting [d, 1] into the following three mesh-intervals:

ree mesh-intervals:
$$\begin{bmatrix} [d, d + \tau] \cup (d + \tau, 1 - \tau] \cup (1 - \tau, 1]. \\
\tau = \min \left\{ \frac{1 - d}{4}, \frac{2\sqrt{\varepsilon}}{\sqrt{\alpha}} \ln N \right\}$$
(8)

Next, on the sub-interval $(d + \tau, 1 - \tau)$, a uniform mesh of N/4 mesh-points is created; on each of the sub-intervals $(d, d + \tau)$ and $(1 - \tau, 1]$ a uniform mesh of N/8 mesh-points is set. It is convenient to take it in practice N = 8n, n > 3.

When $\sigma = \tau = \frac{1}{8N}$ are set to the left, the Shishkin mesh $\overline{\Omega}^N$ transforms into a classical uniform mesh with the transformation parameters σ , τ and a scale N^{-1} from 0 to 1. The following inequalities apply to the mesh Ω^N ,

$$\delta_{h} \leq 2N^{-1} \quad for \qquad 1 \leq h \leq N$$

$$\delta_{h} \geq N^{-1} \quad for \qquad \frac{N}{8} \leq h \leq \frac{3N}{8} \text{ and } \frac{5N}{8} \leq h \leq \frac{7N}{8}$$

$$\delta_{h} \leq N^{-1} \quad for \qquad 1 \leq h \leq \frac{N}{8} \text{ and } \frac{3N}{8} \leq h \leq \frac{N}{2}$$

$$\delta_{h} \leq N^{-1} \quad for \qquad \frac{N}{2} \leq h \leq \frac{5N}{8} \text{ and } \frac{7N}{8} \leq h \leq N$$

$$(10)$$

THE DISCRETE PROBLEM

In this section, a finite element approach with an appropriate Shishkin mesh is used to develop a numerical method for (4).Let for i=1,...,n and $k=1,2,...,N\setminus\{N/2\}$, piecewise linear functional $\operatorname{space} Z_{i,k} \subset H^1_0\big(\Omega^- \cup \Omega^+\big)^n$, that vanishes at $x=\{0,1\}$.This defines the finite element approach for the discrete two-point boundary value problem $Y_{i,k} \in Z_{i,k} \subset H^1_0(\Omega^- \cup \Omega^+)$, for i=1,...,n,

$$\beta_{i}(Y_{i,k}(x), z_{i,k}(x)) = f_{i,k}(z_{i,k})(x) \quad \forall \quad z_{i,k} \in Z_{i,k} \subset H_{0}^{1}(\Omega^{-} \cup \Omega^{+})$$

$$f_{i,k}(z_{i,k})(x) = (f_{i,k}(x), z_{i,k}(x)).$$

$$\beta_{i}(y_{i,k}(x), z_{i,k}(x)) = -\varepsilon \quad (y'_{i,k}(x), z'_{i,k}(x)) + \left(\sum_{i=1}^{n} (a_{ij}(x)y_{j,k}(x)), z_{i,k}(x)\right)$$

$$(11)$$

Lemma suggests that the discrete problem is stable and has a unique solution according to Lax-Migram.





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INTERPOLATION ERROR BOUNDS

Lemma 3: Define $y_{i,k}^*$ as the $Z_{i,k}$ -interpolant of the solution $y_{i,k}$ of (1) on the fitted mesh Ω^N . Then

$$\max_{i=1,\dots,n} \|y_{i,k}^* - y_{i,k}\|_{\Omega^N} \le C(N^{-1}lnN)^2,$$

 \boldsymbol{C} is a constant that is independent of the parameter $\boldsymbol{\varepsilon}~$.

Proof

The solution of this Lemma 3 is obtained by replacing with identical parameters according to the Lemma 5.1 in [13]. **Lemma 4:** Define $y_{i,k}^*$ as the $Z_{i,k}$ interpolant of the solution $z_{i,k}$ of (1) on the fitted mesh Ω^N . Then

$$\max_{i = 1, ..., n} \| y_{i,k}^* - y_{i,k} \|_{\varepsilon} \le C(N^{-1} \ln N)^2,$$

C is a constant that is independent of the parameter ε .

Proof:

The solution of this Lemma 4 is obtained by replacing with identical parameters according to the Lemma 5.2 in [13]. **Lemma 5:** Define $y_{i,k}^*$ as the $Z_{i,k}$ -interpolant of the solution $y_{i,k}$ of (1) on the fitted mesh Ω^N . Then

$$\max_{i=1,...,n} \| y_{i,k}^* - y_{i,k} \|_{\varepsilon,\overline{\Omega}^N} \le C (N^{-1} \ln N)^2.$$

C is a constant that is independent of the parameter.

Proof:

The solution of this Lemma 5 is obtained by replacing with identical parameters according to the Lemma 5.3 in [13].

INTERPOLATION ERROR ESTIMATE AND DISCRETIZATION ERROR

Lemma 6: Define $y_{i,k}$ as the solution of (1) and $Y_{i,k}$ the solution of (4) Suppose that $Z_{i,k} \subset H_0^1(\Omega^- \cup \Omega^{+^N})$. Then $\max_{i = 1, \ldots, n} \left| \beta_i (Y_{i,k} - y_{i,k}, v_i) \right| \le C (N^{-1} \ln N)^2 \quad \| \ z_{i,k} \ \|_{l^2(\Omega^{-N})},$

C is a constant that is independent of the parameter ε

Proof:

The solution of this Lemma 6 is obtained by replacing with identical parameters according to the Lemma 6.1 in [13].

Lemma 7: Define $y_{i,k}^*$ as the $Z_{i,k}$ -interpolant of the solution $y_{i,k}$ of (1) and $Y_{i,k}$ the solution of (4). Then

$$\max_{i=1,\ldots,n}\|Y_{i,k}-y_{i,k}^*\|_{\varepsilon,\overline{\Omega}^N}\leq C(N^{-1}\ln N)^2,$$

C is a constant that is independent of the parameter ε .

Proof: Since $Y_{i,k} - y_{i,k}^* \in Z_{i,k}$,

$$\begin{split} & \parallel Y_{i,k} - y_{i,k}^* \parallel_{\varepsilon}^2 \sum_{,\overline{\Omega}^N} \leq C \, \beta_i \big(Y_{i,k} - y_{i,k}^*, Y_{i,k} - y_{i,k}^* \big) \\ & \leq C \big[\beta_i \big(Y_{i,k} - y_{i,k}, Y_{i,k} - y_{i,k}^* \big) + \beta_i \big(y_{i,k} - y_{i,k}^*, Y_{i,k} - y_{i,k}^* \big) \big] \end{split}$$

Using Lemma 4, with $z_i = Y_{i,k} - y_{i,k}^*$, then gives

$$\|Y_{i,k} - y_{i,k}^*\|_{\varepsilon,\overline{\Omega}^N}^2 \le C (N^{-1} \ln N)^2 \|Y_{i,k} - y_{i,k}^*\|_{\varepsilon,\overline{\Omega}^N}.$$

Cancelling the common factor gives

$$\| Y_{i,k} - y_{i,k}^* \|_{\varepsilon, \overline{\Omega}^N} \le C(N^{-1} \ln N)^2$$

as required.

Theorem 1: Define $y_{i,k}$ as a solution of (1) and $Y_{i,k}$ as a solution of (4). Then

$$\max_{i=1,\ldots,n} \|Y_{i,k} - y_{i,k}\|_{\varepsilon,\overline{\Omega}^N} \le C(N^{-1}\ln N)^2,$$

C is a constant that is independent of the parameter ε .





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Proof: Since

$$\parallel Y_{i,k}-y_{i,k}\parallel_{\varepsilon_{-,\Omega}N}\leq \parallel Y_{i,k}-y_{i,k}^*\parallel_{\varepsilon_{-}\overline{\Omega}^N}+\parallel y_{i,k}^*-y_{i,k}\parallel_{\varepsilon_{-}\overline{\Omega}^N},$$

the result follows by combining Lemmas (3) and (7).

Theorem 2: Let $y_{i,k}$ be the solution of (1) and $Y_{i,k}$ the solution of (5). Then the following parameter uniform error estimate holds

$$\max_{i = 1, \ldots, n} \| Y_{i,k} - y_{i,k} \|_{\varepsilon, \overline{\Omega}^N} \le C(N^{-1} \ln N)^2$$

C is a constant that is independent of the parameter ε

Proof:

The solution of this Theorem is obtained by replacing with identical parameters according to the Theorem 7.3in [13].

NUMERICAL ILLUSTRATIONS

Example: Consider the BVP

$$-E\vec{u}''(x) + A(x)\vec{u}(x) = \vec{f}(x), \ for \ x \in (0,1), \ \vec{u}(0) = \vec{0}, \ \vec{u}(1) = \vec{0} \text{ where } E = diag \ (\varepsilon \ , \varepsilon \), \ A = \begin{pmatrix} 5 & -1 \\ -1 & 5(x+1) \end{pmatrix}, \ \vec{f} = (0,1), \ \vec{u}(0) = \vec{0}, \ \vec{u}(1) = \vec{0} \text{ where } E = (0,1), \ \vec{u}(0) = \vec{0}, \ \vec{u}(1) = \vec{0} \text{ where } E = (0,1), \ \vec{u}(0) = \vec{0}, \ \vec{u}(1) = \vec{0} \text{ where } E = (0,1), \ \vec{u}(0) = \vec{0}, \ \vec{u}(1) = \vec{0} \text{ where } E = (0,1), \ \vec{u}(0) = \vec{0}, \ \vec{u}(1) = \vec{0} \text{ where } E = (0,1), \ \vec{u}(0) = \vec{0}, \ \vec{u}(1) = \vec{0} \text{ where } E = (0,1), \ \vec{u}(0) = \vec{0}, \ \vec{u}(1) = \vec{0} \text{ where } E = (0,1), \ \vec{u}(0) = \vec{0}, \ \vec{u}(1) = \vec{0} \text{ where } E = (0,1), \ \vec{u}(0) = \vec{0}, \ \vec{u}(1) = \vec{0} \text{ where } E = (0,1), \ \vec{u}(0) = \vec{0}, \ \vec{u}(1) = \vec{0} \text{ where } E = (0,1), \ \vec{u}(0) = \vec{0}, \ \vec{u}(1) = \vec{0} \text{ where } E = (0,1), \ \vec{u}(0) = \vec{0}, \ \vec{u}(1) = \vec{0} \text{ where } E = (0,1), \ \vec{u}(0) = \vec{0}, \ \vec{u}(1) = \vec{0} \text{ where } E = (0,1), \ \vec{u}(0) = \vec{0}, \ \vec{u}(1) = \vec{0} \text{ where } E = (0,1), \ \vec{u}(0) = \vec{0}, \ \vec{u}(1) = \vec{0} \text{ where } E = (0,1), \ \vec{u}(0) = \vec{0}, \ \vec{u}(1) = \vec{0} \text{ where } E = (0,1), \ \vec{u}(0) = \vec{0}, \ \vec{u}(1) = \vec{0} \text{ where } E = (0,1), \ \vec{u}(1) = \vec{0} \text{ where } E = (0,1), \ \vec{u}(1) = \vec{0} \text{ where } E = (0,1), \ \vec{u}(1) = \vec{0} \text{ where } E = (0,1), \ \vec{u}(1) = \vec{0} \text{ where } E = (0,1), \ \vec{u}(1) = \vec{0} \text{ where } E = (0,1), \ \vec{u}(1) = \vec{0} \text{ where } E = (0,1), \ \vec{u}(1) = \vec{0} \text{ where } E = (0,1), \ \vec{u}(1) = \vec{0} \text{ where } E = (0,1), \ \vec{u}(1) = \vec{0} \text{ where } E = (0,1), \ \vec{u}(1) = \vec{0} \text{ where } E = (0,1), \ \vec{u}(1) = \vec{0} \text{ where } E = (0,1), \ \vec{u}(1) = \vec{0} \text{ where } E = (0,1), \ \vec{u}(1) = \vec{0} \text{ where } E = (0,1), \ \vec{u}(1) = ($$

 $(1+x^2,2)^T$. For various values of ϵ , N=8k, $k=2^r$, $r=3,\cdots,8$, and $\alpha=1.9$.Based on the general methods outlined in Miller et al. [6], we applied the fitted mesh method to Example to compute the ϵ uniform order of convergence and the ϵ uniform error constant. The results of our computations are summarized in Table 1, which provides the following conclusions:

CONCLUSION

The research presented in this article builds upon the foundational concept pioneered by Miller et al. [6], who focused on convection-diffusion problems. In our paper, we extend this framework by establishing second order parameter uniform convergence for systems of n second order differential equations of reaction diffusion type with same perturbation parameter, incorporating discontinuous source terms. This advancement in methodology allows for a more robust treatment of complex systems, where discontinuities in source terms pose significant challenges. By achieving second order parameter uniform convergence, we enhance the accuracy and reliability of numerical solutions for such systems. Furthermore, our proposed method is not limited to one dimensional problems; it can be extended to tackle higher-dimensional scenarios. This scalability is crucial for addressing real world problems that often involve complex geometries and multi-dimensional phenomena.

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Table 1: Values of D_{ε}^{N} , D^{N} , p^{N} , p^{*} and $C_{p^{*}}^{N}$ for $\varepsilon_{1} = \varepsilon_{2} = \frac{\eta}{64}$.

η	Number of mesh points N				
	64	128	256	512	1024
20	0.3142E-02	0.1550E-02	0.7695E-03	0.3833E-03	0.1913E-03
2-1	0.4518E-02	0.2223E-02	0.1111E-02	0.5593E-03	0.2765E-03
2-2	0.6386E-02	0.3168E-02	0.1577E-02	0.7869E-03	0.3929E-03
2-3	0.8985E-02	0.4474E-02	0.2229E-02	0.1112E-02	0.5558E-03
2-4	0.1260E-01	0.6311E-02	0.3150E-02	0.1573E-02	0.7860E-03
2-5	0.1756E-01	0.8892E-02	0.4449E-03	0.2223E-02	0.1111E-02
2-6	0.1756E-01	0.8892E-02	0.4449E-03	0.2223E-02	0.1111E-02
2-7	0.1756E-01	0.8892E-02	0.4449E-03	0.2223E-02	0.1111E-02
D^N	0.1756E-01	0.8892E-02	0.4449E-03	0.2223E-02	0.1111E-02
P^N	0.9758E+00	0.9952E+00	0.1253E+01	0.1473E+01	
C_P^N	0.9633E+00	0.9453E+00	0.8898E+00	0.7031E+00	0.5032E+00
Computed order of $\vec{\varepsilon}$ uniform convergence, $p^* = 1.293$					
Computed $\vec{\varepsilon}$ -uniform error constant, $C_{p^*}^N = 0.8233$					





RESEARCH ARTICLE

Study on Awareness about Recent Regulations in UPI among Users

R.Ragavi¹ and PL. Meenakshi²

¹Post Graduate Student, Department of Mathematics, Avinashilingam Institute for Home Science and Higher Education for Women, Tamil Nadu, India.

²Assistant Professor, Department of Mathematics, Avinashilingam Institute for Home Science and Higher Education for Women, Tamil Nadu, India.

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*Address for Correspondence

R.Ragavi

Post Graduate Student, Department of Mathematics,

Avinashilingam Institute for Home Science and Higher Education for Women, Tamil Nadu, India.

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ABSTRACT

Unified Payment Interface (UPI) is the most used digital payment method that keeps growing every year with new users due to its ease UPI is a secure and convenient way to make payments without carrying physical money. A recent study is focused on analyzing the awareness of the regulations set by NPCI (National Payment Corporation of India) that limit the number of transactions to 20 per 24 hours and the maximum payable amount to 1 lakh, cumulatively. The study was carried out with the primary data collected by simple random sampling with a sample size of 69 users. The study analyses the level of awareness among users about interchange fees and limits. Some statistical tools used are the chi-square test and percentage analysis. The analysis pointed out that around 51% of the users were aware of the number of transactions in a day and around 61% of the people had known the highest amount that can be sent.

Keywords: Digital Payment, UPI, NPCI, Awareness, Transactions.

INTRODUCTION

Unified Payment Interface (UPI) is one of the fastest-growing digital transaction methods that has gained prominence in recent years. People are getting more and more dependent on digital technologies, and many people prefer digital ways of payment then cash transactions due to its ease. Unified Payment Interface was developed by the National Payment Corporation of India (NPCI) in the year 2016, NPCI is an organization that settles retail payment in India. It is introduced by the Reserve Bank of India (RBI) and the Indian Banks Association (IBA). UPI enables the user to send and receive money from one bank account to another, facilitating person-to-person and person-to-merchant transactions. It is a real-time payment system, meaning the money will be exchanged





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immediately without delay. It enables the user to send and receive money with unique UPI IDs which use VPA (Virtual Payment Address) to transfer the money securely to other bank accounts linked to that UPI ID. Since VPA is unique no fake ID can be created which makes it highly secure. It provides various banking features to the users like balance checking and amount transfer which reduces the need to visit banks or ATMs.

STATEMENT OF THE PROBLEM

As the economy advances, online transactions are gaining popularity rapidly. More and more individuals are opting for online payment methods like

- 1. UPI,
- 2. digital currencies,
- 3. cryptocurrencies and Internet banking

instead of cash due to their convenience. UPI transactions amount to **62**% of digital payment transactions in FY 2022-23. The study aims to investigate the potential impact of recent NPCI regulations on UPI usage.

OBJECTIVE OF THE STUDY

The objective of this study is to examine the level of awareness regarding recent limitations on the number of transactions that can be initiated in UPI and the maximum amount that can be transferred through UPI apps.

RESEARCH METHODOLOGY

A descriptive research design has been employed to analyze the awareness among UPI users. Random sampling method was used to take the sample and the sample size is 69. The data were collected from students. Structured methods of questionnaires have been employed to collect primary data from the respondents.

REVIEW OF LITERATURE

Dr.D. Durairaj & Princy Joseph (2019) UPI is more feasible than the mobile wallet if server-related issues are improved and if the right awareness about it is produced for young minds. It should attract the illiterate people of the society to use UPI to make it more inclusive. Rewarding UPI payments would sufficiently contribute to the growth of UPI. Amogh Das & Ashish Das (2022) Year-on-year growth and UPI transactions on a daily basis have improved and the number of transactions done from person to person and through person to merchant has increased significantly. Due to the government's intention to support UPI payments banks received monetary support. Putta Sai Kumar (2022) The level of satisfaction of customers with UPI is independent of the number of transactions done and the payments. The satisfaction level is not influenced by the complaints, issues, and transaction services provided by the banks.

Statistical Methods Used for Analysis

SPSS software was employed to analyze the data in the following methods.

- 1. Percentage Analysis
- 2. Chi-Square test
- 3. ANOVA table.

PERCENTAGE ANALYSIS

DISTRIBUTION OF UPI AWARENESS BASED ON AGE

Null hypothesis H_0 : There is no significant relation between age and the level of awareness. Alternative hypothesis H_1 : There is a significant relation between age and the respondents' awareness level. $\alpha = 0.05$ level of significance.





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Interpretation

Number of transactions limit

P= 0.879 is greater than 0.05 hence null hypothesis is accepted. There is no significant relation between number of transaction limits and age.

Payment amount limit

P= 0.053 is higher than 0.05 hence null hypothesis is accepted. There is no significant relation between payment amount limit and age.

Prepaid Instruments

P = 0.597 is higher than 0.05, hence the null hypothesis is accepted and there is no significant relation between the level of awareness of prepaid instruments and age.

DISTRIBUTION OF AWARENESS ABOUT THE HIGHEST AMOUNT THAT CAN BE TRANSFERRED:

Null Hypothesis H₀: There is no significant relation between the number of transactions done by a person and the awareness

Alternative hypothesis H₁: There is a significant relationship between the number of transactions done by a person and awareness.

 α = 0.05 has been considered as the level of significance.

Chi-Square Test

Interpretation

P value = 0.320

P value is greater than 0.05, hence null hypothesis is accepted.

There is no significant relation between some transactions done by individuals and the level of awareness about the highest amount limit.

FINDINGS OF THE SURVEY CARRIED OUT AMONG UPI USERS

94% of the people surveyed used UPI apps.

50.7% of the people surveyed used LESS than 5 transactions in a week.

20.9% of the people surveyed used LESS than 10 transactions in a week.

13.4% of the people surveyed used LESS than 15 transactions in a week.

14.9% of the people surveyed used MORE than 15 transactions in a week.

50.7% of the people surveyed were aware of the highest number of transactions that can be done through UPI in a day.

53.4% of the people surveyed used UPI transactions for merchant transfers of money.

60.9% of the people knew the highest limit that could be sent through UPI.

53.6% of the people surveyed have been denied to pay through UPI and requested to pay through other means at least once

68.1 % of the people have used prepaid instruments.

5.8% of the users have encountered UPI fraud or lost money through UPI payments.

LIMITATIONS OF THE STUDY

- 1. The smaller size of the sample (69).
- 2. Less Geographical area covered.
- 3. The survey is limited only to the literates.





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CONCLUSION

The recent regulations have not affected the growth of UPI payments. Most of the persons surveyed used UPI payments for merchant transfer thus helping to formalize the economy. 4% of the people surveyed have encountered UPI frauds thus with digitalizing the economy we have to increase online security and compliance.

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RESEARCH ARTICLE

Detecting and Mitigating Cybersecurity Threats in Smart Grids using **KMeans Clustering**

Ramya Hyacinth Lourdusamy¹ and Gomathi Venugopal²

¹Department of Electrical and Electronics Engineering, Loyola-ICAM College of Engineering and Technology, Chennai, India.

²Department of Electrical and Electronics Engineering, College of Engineering, Guindy, Anna University, Chennai, India.

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*Address for Correspondence Ramya Hyacinth Lourdusamy

Department of Electrical and Electronics Engineering, Loyola-ICAM College of Engineering and Technology, Chennai, India.

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ABSTRACT

In contemporary years, the integration of advanced digital meters, smart meters, and Phasor Measurement Units (PMUs) into power system networks has revolutionized real-time monitoring capabilities. These metering infrastructures are indispensable for facilitating decision-making processes and ensuring the smooth operation of power systems. However, their crucial role also renders them vulnerable to emerging threats, particularly in the realm of cybersecurity. One of the most pressing threats facing power system metering infrastructure is the occurrence of False Data Injection Attacks (FDIAs). These malicious attacks, resulting from the digital transformation of power systems, entail the injection of falsified data into the metering infrastructure. Such anomalies can severely compromise the reliability and stability of power system networks, undermining their integrity and posing significant operational risks. Traditional approaches to enhancing data reliability, such as statistical methods, have proven inadequate in effectively identifying and mitigating FDIAs. Consequently, there is a pressing need to leverage more advanced techniques, such as Machine Learning (ML), to combat this evolving threat landscape. This paper addresses the challenge of detecting FDIAs using ML algorithms and specifically, tailored to detect and mitigate FDIAs in power system networks. To evaluate the efficacy of our approach, the algorithm is tested using a dataset comprising real-time PMU data from the Australian Grid, artificially injected with false data. Additionally, the performance of the model is assessed in terms of precision and recall, providing valuable insights into its effectiveness in securing power system metering infrastructure against FDIAs.

Keywords: One of the most pressing threats facing power system metering infrastructure is the occurrence of False Data Injection Attacks (FDIAs).





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INTRODUCTION

Data-driven systems [1] play a pivotal role in modern society, facilitating critical operations across various domains such as energy management, transportation, finance, and healthcare. These systems rely heavily on the integrity and accuracy of the data they process to make informed decisions and ensure reliable performance [2]. However, the proliferation of interconnected devices and the growing complexity of data networks have introduced new vulnerabilities, leaving these systems susceptible to malicious attacks. One such threat is the FDIA, a sophisticated form of cyberattack intended to undermine the integrity of data within a system[3]. In a false data injection attack, adversaries clandestinely inject fabricated or manipulated data into the system to deceive its operations and undermine its functionality. These attacks pose a significant threat to the reliability, security, and safety of datadriven systems, with potentially severe consequences ranging from financial losses [4] to operational disruptions [5] and even threats to public safety. Power systems equipped with advanced Information and Communication Technology are being transformed into a smart grid. Smart Grids are designed to enhance the reliability and efficiency of energy distribution [6]. Integrating and analysing data from these multiple modalities enables utilities and operators to enhance situational awareness, optimize operations, improve reliability, and support decisionmaking processes in power systems. The multimodal data collected from Smart meters, Phasor Measurement Units (PMUs) and other Advanced Metering Infrastructure (AMI) serves as the foundation for decision-making processes and control actions [7]. However, FDIAs can introduce inaccuracies into the data collected from sensors and meters, leading to incorrect load forecasting, faulty control decisions, and ultimately, disruptions in energy supply.

The impact of false data injection attacks can be particularly devastating in critical infrastructure sectors such as smart grids, where accurate data is essential for efficient energy distribution and grid stability [8]. In a smart grid context, for example, attackers may inject false data into sensor readings to manipulate load forecasts, disrupt energy distribution, or even cause widespread blackouts. Detecting and mitigating false data injection attacks present significant challenges due to the stealthy nature of these attacks and the complexity of data-driven systems. Traditional security measures such as firewalls and encryption may not be sufficient to defend against such threats, as attackers can exploit vulnerabilities in data processing algorithms and protocols to evade detection. Therefore, there is an urgent need for advanced anomaly detection techniques that can effectively identify and thwart false data injection attacks in real time. By detecting and mitigating FDIAs, the reliability of energy supply in Smart Grids can be safeguarded, ensuring uninterrupted power delivery to consumers [9]. Also, FDIAs can inject false data into Phasor Measurement Units (PMUs) and other monitoring devices, leading to misleading information about the state of the grid. This can undermine the stability of the Smart Grid, potentially causing voltage instability, frequency deviations, and even cascading failures [10]. This enhances the resilience of Smart Grids to cyber threats. With the increasing digitization of power systems, cybersecurity data becomes crucial for monitoring and protecting against cyber threats, such as malware, intrusion attempts, and denial-of-service attacks. The use of KMeans clustering-a popular unsupervised machine learning algorithm—for detecting false data injection attacks in data-driven systems is focussed in this article. By leveraging the inherent structure of the data and identifying deviations from normal patterns, a robust and efficient approach for safeguarding the integrity of data in critical infrastructure and other mission-critical applications is developed. Through empirical evaluation and analysis, the effectiveness and practical applicability of the proposed methodology in mitigating the threat posed by false data injection attacks are demonstrated.

REVIEW OF EXISTING LITERATURE

FDIAs involve injecting falsified data into sensors or measurement devices to manipulate system behaviour. Studies have investigated the impact of FDIAs on different types of systems, including power grids, water distribution networks, and industrial control systems. Research has focused on developing detection and mitigation techniques to counter FDIAs, ranging from statistical anomaly detection methods to machine learning-based approaches. Anomaly detection is a crucial component of cybersecurity, aiming to identify abnormal patterns or behaviours





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indicative of malicious activity [11]. Traditional anomaly detection techniques include statistical methods such as mean-variance analysis, time-series analysis, and clustering-based approaches [12]. Statistical methods may struggle to distinguish between genuine anomalies and normal variations in system behaviour, leading to a high false positive rate [13]. These approaches often rely on predefined thresholds or assumptions about the distribution of data, which may not hold true in complex and dynamic systems. Statistical techniques may not be effective for detecting subtle and sophisticated attacks, such as those involving gradual changes in data over time.

More recent advancements in anomaly detection involve Machine Learning (ML) and data mining techniques, including supervised, unsupervised, and semi-supervised learning algorithms. ML models can learn patterns and relationships in data to identify anomalies. Common ML algorithms used for anomaly detection include support vector machines (SVMs), k-nearest neighbours (KNN), isolation forests, and autoencoders [14]. ML algorithms require labelled training data for supervised learning, which may be scarce or expensive to obtain, particularly for rare and complex anomalies. Unsupervised and semi-supervised ML algorithms may be used for problems involving unlabelled real-time data. ML models could be vulnerable to adversarial attacks [15], where attackers manipulate input data to circumvent detection by the anomaly detection system. Research has explored the application of deep learning models, such as autoencoders and recurrent neural networks, for detecting complex anomalies in high-dimensional data streams [16]. KMeans clustering, a popular unsupervised learning algorithm, has found various applications in cybersecurity, particularly in anomaly detection and intrusion detection systems [17]. Despite significant research efforts in detecting FDIAs, several gaps remain in the literature. These gaps indicate areas requiring additional investigation and development to bolster the effectiveness and robustness of FDIAs detection.

Some of the key gaps include

- Dynamic and Adaptive Detection: Existing detection techniques often rely on static thresholds or predefined
 models to identify anomalies. However, FDIAs can be highly adaptive and stealthy, evolving to evade
 detection. There is a need for dynamic and adaptive detection methods that can continuously learn and
 evolve in response to changing attack strategies.
- Scalability and Efficiency: Numerous detection algorithms encounter scalability challenges when utilized inlarge-scale systems with high-dimensional data streams, such as Smart Grids. Scalability becomes crucial for real-time detection in such scenarios. There is a need for scalable detection techniques that can handle the volume, velocity, and variety of data generated by modern cyber-physical systems.
- Multimodal Data Fusion: Smart Grids are complex electrical infrastructures that collect data from different
 meters not limited to Supervisory Control and Data Acquisition (SCADA), PMU, and Smart Meters. These
 meters have different sampling rates and capture parameters of different kinds. Thus, data obtained in realtime is multimodal data. Integrating information from multiple data sources can enhance the detection
 accuracy and resilience to attacks. However, there is limited research on effective techniques for multimodal
 data fusion in the context of FDIA detection.
- Adversarial Robustness: FDIA detection systems may be susceptible to adversarial attacks, where attackers tactically engineer input data to evade detection. Research is needed to develop detection techniques that are robust to adversarial perturbations and can withstand sophisticated evasion strategies employed by attackers.
- Real-world Evaluation: While numerous detection algorithms have been proposed in the literature, few have been extensively evaluated in real-world settings or deployed in operational environments. There is a lack of empirical studies that assess the performance, scalability, and practical utility of FDIAs detection techniques under realistic conditions.
- Cross-domain Adaptability: FDIAs detection methods developed for specific domains, such as power systems or industrial control systems, may not be directly applicable to other domains due to differences in data characteristics and system architectures. There is a need for cross-domain adaptable detection techniques that can be effectively deployed across diverse cyber-physical systems.





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On the whole, previous research in these areas and the listed research gaps have laid the groundwork for developing more robust and effective cybersecurity solutions. By leveraging insights from past studies, detecting and mitigating cybersecurity threats including FDIAs using KMeans clustering is presented.

METHODOLOGY

The most popularly applied concept in distance-based ML approaches is the Euclidean distance. These values act as a tool for identifying malicious data. KMeans clustering helps to group a dataset with N entities into K -clusters. Clusters are formed by assigning data to the closest centroid by calculating distances. Thus, by calculating the distance between the data and the centroid, the ones with the largest distances are identified as anomalous data. In this case, the data is grouped into binary clusters of normal/ anomalous data. The mathematical formulation is explained below: If the dataset is represented as $dt_1, dt_2, ..., dt_n$, then they are grouped into k clusters $C = \{C_1, C_2, ..., C_k\}$. The means of these clusters be represented as $\mu_1, \mu_2, ..., \mu_k$; where μ_i is the mean of points in C_i . Normally, K -Means clustering follows two steps iteratively. They are the assignment step and the update step [18].

Assignment step

Every data point dt_p is assigned to a cluster with the closest mean. This is achieved by calculating the least square Euclidean distance.

$$C_i^{(t)} = \left\{ dt_p : \|dt_p - \mu_i^{(t)}\|^2 \le \|dt_p - \mu_j^{(t)}\|^2 \right\} \forall j, 1 \le j \le k \tag{1}$$

Thus, every data point dt_n is assigned only to one cluster C_i in the assignment step.

Update step

The centroid or mean of the cluster to which the data point dt_p was assigned in the previous step is recalculated and updated.

$$\mu_i^{(t+1)} = \frac{1}{|C_i^{(t)}|} \sum_{dt_j \in C_i^{(t)}} dt_j \tag{2}$$

The above two steps are repeated iteratively till the recalculated mean no longer changes. The data point that has the largest Euclidean distance is identified as the malicious data.

RESULTS AND DISCUSSION

Experimental Setup

Real-Time data obtained from 15 Phasor Measurement Units (PMUs) deployed at three different time zones in Australia that are sampled at three different sampling rates 30, 60 and 120 Hz at different offsets is considered for evaluating the methodology. Initially, the data in three different time zones with different offsets are aligned and the missing and duplicate data are removed. To ensure the availability of data at uniform intervals, missing values are interpolated using curve fitting techniques. Also, the measurements of PMU include voltage magnitudes, voltage angles and frequencies that are at different scales. Robust scaling is performed to prevent features with larger scales from dominating the analysis. In addition, dimensionality reduction using Principal Component Analysis (PCA) is performed as high-dimensional time series data test data is considered for analysis.

Let H denotes the Jacobian matrix obtained from the system configuration. Under conditions, when attack vector a is composed of a linear combination of the column vectors within H. That is a false data injection attack vector, a = Hc, is designed and injected along with the measurement vector, $\Box_{\Box} \coloneqq \Box + \Box$. This intruder-attacked measurement surreptitiously escapes the bad data detection tests (Chi-square and Largest Normalised Residual Tests). For the sake of analysis, 5% of the measurement is intentionally corrupted.





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Findings and Interpretations

The elbow method plot for the considered dataset is given in Figure 1. This demonstrates the correlation between the number of clusters (K) and the within-cluster sum of squares (WCSS). The elbow method aids in identifying the optimal number of clusters by pinpointing the juncture where the rate of decrease in WCSS begins to decelerate, signifying diminishing returns from further cluster additions. From Figure 1, it is evident that K for the considered data set could be taken as two or three. On applying K = 2, the scatter plot obtained is shown in Figure 2 and that obtained by considering K = 3 is shown in Figure 3. Outliers, which are observed away from the clustered data points on the scatterplot, are recognized as anomalous data and are excluded before being utilized as inputs for other decision-making operations related to power systems.

CONCLUSION

KMeans clustering contributes to addressing several of the gaps in detecting false data injection attacks (FDIAs) by leveraging its capabilities in unsupervised learning and pattern recognition. From the analysis, it is evident that KMeans Clustering is capable of:

- Dynamic and Adaptive Detection: KMeans can be used to dynamically cluster data points based on their similarity, allowing for the detection of evolving patterns and anomalies. By periodically re-training the KMeans model with new data, it can adapt to changes in system behaviour and identify emerging attack patterns.
- Scalability and Efficiency: KMeans is known for its scalability and efficiency, particularly when dealing with large-scale datasets. By efficiently partitioning data into clusters, KMeans helps to reduce the computational complexity of anomaly detection algorithms, enabling real-time detection in high-dimensional and highvolume data streams.
- Multimodal Data Fusion: KMeans clustering is applied to integrate information from multiple data modalities
 by clustering data points based on their combined features. By clustering multimodal data streams, KMeans
 can identify complex patterns and correlations that may indicate anomalous behaviour across different data
 sources.
- Adversarial Robustness: KMeans clustering is inherently robust to small perturbations in data, making it
 resistant to simple adversarial attacks. Additionally, ensemble approaches combining multiple KMeans
 models or incorporating robust clustering techniques can further enhance the resilience of FDIA detection
 systems against adversarial manipulation.
- Privacy Preservation: KMeans clustering can be applied to analyse data in a privacy-preserving manner by
 clustering data points based on their similarities while preserving individual privacy. Differential privacy
 techniques can be incorporated to ensure that sensitive information is not revealed during the clustering
 process.
- Real-world Evaluation: KMeans clustering provides a transparent and interpretable framework for analysing
 data and identifying clusters, making it suitable for real-world evaluation of FDIAs detection techniques. By
 applying KMeans to real-world datasets, researchers can assess the performance and practical utility of
 detection algorithms under realistic conditions.
- Cross-domain Adaptability: KMeans clustering is a domain-agnostic technique that can be applied across
 diverse cyber-physical systems and domains. By adapting the feature representation and clustering
 parameters, KMeans can be effectively deployed in different environments to detect FDIAs across various
 systems.
- In summary, KMeans clustering offers a versatile and scalable approach to addressing several of the gaps in FDIA detection. By leveraging its capabilities in unsupervised learning and pattern recognition, KMeans can contribute to the development of robust and efficient detection techniques for identifying and mitigating FDIAs in complex and dynamic cyber-physical systems.





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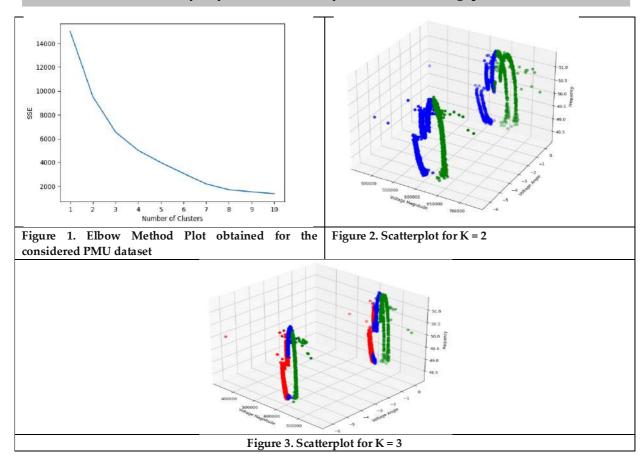
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RESEARCH ARTICLE

Pair Mean Cordial Labeling of Hexagonal Snake, Irregular Quadrilateral Snake and Triple Triangular Snake

R. Ponraj¹ and S. Prabhu²

¹Department of Mathematics, Sri Paramakalyani College, Alwarkurichi-627 412, India.

²Research Scholar, Reg. No.: 21121232091003, Sri Paramakalyani College, Alwarkurichi-627 412, India.

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*Address for Correspondence R. Ponraj

Department of Mathematics, Sri Paramakalyani College, Alwarkurichi-627 412, India. Email: ponrajmaths@gmail.com



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ABSTRACT

Consider a (p,q) graph G = (V,E). Specify $\varrho = \begin{cases} \frac{p}{2} & even \ p \\ \frac{p-1}{2} & odd \ p, \end{cases}$ and $\Upsilon = \{\pm 1, \pm 2, \dots, \pm \varrho\}$ referred to as the

label set. Let's consider a mapping $\lambda: V \to \Upsilon$ where, for every even p, distinct labels are assigned to the various elements of V in Υ , and for every odd p, distinct labels are assigned to the p-1 vertices of V in Υ , with a label that is repeated for the lone vertex. After that φ is referred to as a pair mean cordial labeling (PMC-labeling) if regarding every $\mu\nu$ edge within G, there is a label for $\frac{\varphi(\mu)+\varphi(\nu)}{2}$ if $\varphi(\mu)+\varphi(\nu)$ is even and $\frac{\varphi(\mu)+\varphi(\nu)+1}{2}$ if $\varphi(\mu)+\varphi(\nu)$ is odd ensue that $\left|\overline{\mathbb{S}}_{\varphi_1}-\overline{\mathbb{S}}_{\varphi_1^c}\right|\leq 1$ in which $\overline{\mathbb{S}}_{\varphi_1}$ are signify the quantity of edges designated with 1 and the quantity of edges that are unlabeled with 1 in that order. A pair mean cordial graph (PMC-graph) is defined as a graph G with PMC-labeling. In our study, we examine the PMC-labeling of several snake graphs.

Keywords: hexagonal snake, irregular quadrilateral snake, comb triangular graph, quadrilateral snake and triple triangular snake

INTRODUCTION

In our study, we consider a graph as a undirected, finite graph that has no looping and only one edge. Graph labeling is a well-known field of study in graph theory, with an extensive collection of literature and unresolved issues covering a wide range of graph kinds. Rosa[17] initially put up the graph labeling idea in the 1960s. We adhere to F. Harary [4] for different graphs, theoretical notations, and nomenclature and Gallian [3] provides a thorough analysis of different graph labeling techniques.
 I. Cahit [2] suggested the idea of cordial graphs and looking at the





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cordial related graphs in [1,5-8,13-16,18-20]. Labeling graphs, and more specifically, cordial labeling graphs, was a topic of considerable interest. PMC-labeling is a notion that was first proposed to us in [9] and in [10-12], we looked at the pair mean cordiality of several family graphs. We examine PMC-labeling of several snake graphs, such as the hexagonal snake, irregular quadrilateral snake, comb triangular graph, double quadrilateral snake, alternate double quadrilateral snake, triple triangular snake and triple quadrilateral snake, in this article.

Preliminaries

Definition 2.1. The hexagonal snake HS_r is a graph that is created by taking a path $\mu_1\mu_2, ..., \mu_r$ of length r by adding μ_i and μ_{i+1} to three new vertices ν_i , η_i and ω_i , $1 \le i \le r-1$.

Definition 2.2. A graph known as the irregular quadrilateral snake IQ_r is produced by following a path with vertex set $V(IQ_r) = \{\mu_i, \nu_i, \omega_i \mid 1 \le i \le r \text{ and } 1 \le j \le r - 2\}$ and edge set $E(IQ_r) = \{\mu_i \nu_i, \nu_i \omega_i, \omega_i \mu_{i+2} : 1 \le i \le r - 2\} \cup \{\mu_i \mu_{i+1} : 1 \le i \le r - 1\}.$

Definition 2.3. The double quadrilateral graph TQ_r is a graph that is created by taking a path $\mu_1\mu_2,...\mu_r$ of length r by adding μ_i and μ_{i+1} to new vertices ν_i , ζ_i , η_i and ω_i respectively and then adding ν_i and ζ_i , η_i and ω_i , $1 \le i \le r-1$.

Definition 2.4. A graph known as the alternate double quadrilateral snake $A(DQ_r)$ is produced by joining two alternative quadrilateral snakes which share a path.

Definition 2.5. The triple triangular graph TT_r is made by appending μ_i and μ_{i+1} to new vertices ν_i, ζ_i, η_i and ω_i , $1 \le i \le r - 1$ on a path of length r.

Definition 2.6. The triple quadrilateral graph TQ_r is a graph that is created by taking a path $\mu_1\mu_2,...\mu_r$ of length r by adding μ_i and μ_{i+1} to new vertices ν_j , η_j , ω_j , ζ_j , ξ_j and δ_j respectively and then adding ν_i and η_i , ω_i and ζ_i , ξ_i and δ_i , $1 \le i \le r - 1$.

3. Main Theorems

Theorem 3.1. [10] The quadrilateral snake Q_r is a PMC-graph for all $r \ge 2$.

Theorem 3.2. [10] The triangular snake T_r is a PMC-graph for all $r \ge 2$.

Theorem 3.3. The hexagonal snake HS_r is a PMC-graph for all $r \ge 2$.

Proof. Let $V(HS_r) = \{\mu_i | 1 \le i \le r+1 \} \cup \{\nu_i, \zeta_i, \eta_i, \omega_i | 1 \le i \le r \}$.

Additionally $E(HS_r) = \{\mu_i \eta_i, \mu_i v_i, \eta_i \omega_i, v_i \zeta_i, \zeta_i \mu_{i+1}, \omega_i \mu_{i+1} | 1 \le i \le r \}$. Certainly, the hexagonal snake HS_r consists of 6r edges and 5r + 1 vertices.

Case (A): For odd r

Allocate the labels $-1, -6, ..., \frac{-5r+3}{2}$ corresponding to $\mu_1, \mu_3, ..., \mu_r$ and $4, 9, ..., \frac{5r-7}{2}$ to $\mu_2, \mu_4, ..., \mu_{r-1}$ in that order. Fix the label 1 with μ_{r+1} . Designate the labels $2, 7, ..., \frac{5r-1}{2}$ according to $v_1, v_3, ..., v_r$ and $-4, -9, ..., \frac{-5r+7}{2}$ to $v_2, v_4, ..., v_{r-1}$ in that order. Consequently, allocate the labels $-2, -7, ..., \frac{-5r+11}{2}$ corresponding to $\zeta_1, \zeta_3, ..., \zeta_{r-2}$ and $5, 10, ..., \frac{5r-5}{2}$ to $\zeta_2, \zeta_4, ..., \zeta_{r-1}$ in that order. Fix the label $\frac{-5r-1}{2}$ with ζ_r . Designate the labels $3, 8, ..., \frac{5r+1}{2}$ according to $\eta_1, \eta_3, ..., \eta_r$ and $-5, -10, ..., \frac{-5r+5}{2}$ to $\eta_2, \eta_4, ..., \eta_{r-1}$ in that order. Eventually, allocate the labels $-3, -8, ..., \frac{-5r+9}{2}$ corresponding to $\omega_1, \omega_3, ..., \omega_{r-2}$ and $6, 11, ..., \frac{5r-3}{2}$ to $\omega_2, \omega_4, ..., \omega_{r-1}$ in that order. Fix $\frac{-5r+1}{2}$ with the vertex ω_r . Subsequently, the edges $\omega_1, \omega_2, \omega_3, ..., \omega_{r-2}$ and $\omega_1, \omega_2, \ldots, \omega_{r-2}$ and ω_1, ω_2 $\mu_i\eta_i,\mu_i\nu_i \text{ for } i=1,3,...,r,\ \nu_i\zeta_i,\eta_i\omega_i, \text{ for } i=2,4,...,r-1,\ \zeta_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-2 \text{ and } \eta_r\omega_r \text{ are designated } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1,\ \lambda_i\mu_{i+1},\omega_i\mu_{i+1}, \text{ for } i=1,3,...,r-1$ with 1 and the remaining edges that are unlabeled with 1.

Case (B): For even r

Now designate the labels $-1, -6, \dots, \frac{-5r+8}{2}$ corresponding to $\mu_1, \mu_3, \dots, \mu_{r-1}$ and $4, 9, \dots, \frac{5r-2}{2}$ to $\mu_2, \mu_4, \dots, \mu_r$ in that order. Fix the label 1 with μ_{r+1} . Allocate the labels 2,7, ..., $\frac{5r-8}{2}$ according to $\nu_1, \nu_3, \dots, \nu_{r-1}$ and $-4, -9, \dots, \frac{-5r+2}{2}$ to $\nu_2, \nu_4, \dots, \nu_r$ in that order. Consequently, designate the labels $-2, -7, \dots, \frac{-5r+8}{2}$ corresponding to $\zeta_1, \zeta_3, \dots, \zeta_{r-1}$ and $5, 10, \dots, \frac{5r}{2}$ to $\zeta_2, \zeta_4, \dots, \zeta_r$ in that order. Allocate the labels $3, 8, \dots, \frac{5r-4}{2}$ according to $\eta_1, \eta_3, \dots, \eta_{r-1}$ and $-5, -10, \dots, \frac{-5r}{2}$ to $\eta_2, \eta_4, \dots, \eta_r$





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in that order. Eventually, designate the labels $-3, -8, \ldots, \frac{-5r+4}{2}$ corresponding to $\omega_1, \omega_3, \ldots, \omega_{r-1}$ and $6, 11, \ldots, \frac{5r-8}{2}$ to $\omega_2, \omega_4, \ldots, \omega_{r-2}$ in that order. Fix 1 with the vertex ω_r . Subsequently, the edges $\mu_i \eta_i, \mu_i \nu_i$, $\zeta_i \mu_{i+1}, \omega_i \mu_{i+1}$ for $1 \le i \le r-1$ and i is odd, $\nu_i \zeta_i$, for $1 \le i \le r$ and i is even, $\eta_i \omega_i$, $1 \le i \le r-2$ and i is even and $\eta_r \omega_{r+1}$ are designated with 1 and the remaining edges that are unlabeled with 1. In each instance, $\overline{\mathbb{S}}_{\varphi_1} = 3r = \overline{\mathbb{S}}_{\varphi_1^c}$.

Theorem 3.4. The irregular quadrilateral snake IQ_r is a PMC-graph for all $r \ge 2$.

Proof. Let $V(IQ_r) = \{\mu_i \mid 1 \le i \le r\} \cup \{\nu_i, \omega_i \mid 1 \le i \le r-2\}$. Additionally $E(IQ_r) = \{\mu_i \nu_i, \nu_i \omega_i, \omega_i \mu_{i+2} \mid 1 \le i \le r-2\} \cup \{\mu_i \mu_{i+1} \mid 1 \le i \le r-1\}$. Certainly, the irregular quadrilateral snake IQ_r consists of 4r-7 edges and 3r-4 vertices.

Case (A): For odd r

Allocate the labels 2,5, ..., $\frac{3r-5}{2}$ corresponding to $\mu_1, \mu_3, ..., \mu_{r-2}$ and $-2, -5, ..., \frac{-3r+5}{2}$ to $\mu_2, \mu_4, ..., \mu_{r-1}$ in that order. Fix the label 1 with μ_r . Designate the labels $-1, -4, ..., \frac{-3r+7}{2}$ according to $v_1, v_3, ..., v_{r-2}$ and $4, 7, ..., \frac{3r-7}{2}$ to $v_2, v_4, ..., v_{r-3}$ in that order. Consequently, allocate the labels $3, 6, ..., \frac{3r-9}{2}$ corresponding to $\omega_1, \omega_3, ..., \omega_{r-4}$ and $-3, -6, ..., \frac{-3r+9}{2}$ to $\omega_2, \omega_4, ..., \omega_{r-3}$ in that order. Fix $\frac{3r-5}{2}$ with the vertex ω_r .

Case (B): For even r

Now designate the labels 2,5, ..., $\frac{3r-8}{2}$ corresponding to $\mu_1, \mu_3, ..., \mu_{r-3}$ and $-2, -5, ..., \frac{-3r+8}{2}$ to $\mu_2, \mu_4, ..., \mu_{r-2}$ in that order. Fix the label 1 with μ_{r-1} and $\frac{-3r+4}{2}$ with μ_r . Allocate the labels $-1, -4, ..., \frac{-3r+10}{2}$ according to $v_1, v_3, ..., v_{r-3}$ and $4,7, ..., \frac{3r-4}{2}$ to $v_2, v_4, ..., v_{r-2}$ in that order. Consequently, designate the labels $3,6, ..., \frac{3r-6}{2}$ corresponding to $\omega_1, \omega_3, ..., \omega_{r-3}$ and $-3, -6, ..., \frac{-3r+9}{2}$ to $\omega_2, \omega_4, ..., \omega_{r-2}$ in that order. In each instance, the edges $\mu_i v_i, \omega_i v_i$ for $1 \le i \le r-2$ are designated with 1 and the remaining edges that are unlabeled with 1. Subsequently, $\overline{\mathbb{S}}_{\varphi_1} = 2r-4$ and $\overline{\mathbb{S}}_{\varphi_1^c} = 2r-3$.

Theorem 3.5. The comb triangular graph CT_r is a PMC-graph for all $r \ge 2$.

Proof. Let $V(CT_r) = \{\mu_i, \nu_i, \omega_j \mid 1 \le i \le r \text{ and } 1 \le j \le r-1 \}$. Additionally $E(CT_r) = \{\mu_i \omega_i, \mu_{i+1} \omega_i, \mu_i \mu_{i+1}, \mu_j \nu_j : 1 \le j \le r \text{ and } 1 \le i \le r-1 \}$. Certainly, comb triangular graph CT_r consists of 4r-3 edges and 3r-1 vertices.

Case (A): For odd r

Allocate the labels $-1, -4, ..., \frac{-3r+1}{2}$ corresponding to $\mu_1, \mu_3, ..., \mu_r$ and $4, 7, ..., \frac{3r-1}{2}$ to $\mu_2, \mu_4, ..., \mu_{r-1}$ in that order. Designate the labels $2, 5, ..., \frac{3r-5}{2}$ according to $\nu_1, \nu_3, ..., \nu_{r-2}$ and $-2, -5, ..., \frac{-3r+5}{2}$ to $\nu_2, \nu_4, ..., \nu_{r-1}$ in that order. Fix the label 1 with ν_r . Consequently, allocate the labels $3, 6, ..., \frac{3r-3}{2}$ corresponding to $\omega_1, \omega_3, ..., \omega_{r-2}$ and $-3, -6, ..., \frac{-3r+3}{2}$ to $\omega_2, \omega_4, ..., \omega_{r-1}$ in that order.

Case (B): For even r

Designate the labels $-1, -4, \dots, \frac{-3r+4}{2}$ corresponding to $\mu_1, \mu_3, \dots, \mu_{r-1}$ and $4,7, \dots, \frac{3r-4}{2}$ to $\mu_2, \mu_4, \dots, \mu_{r-2}$ in that order. Fix the label 1 with μ_r . Allocate the labels $2,5,\dots,\frac{3r-2}{2}$ according to v_1,v_3,\dots,v_{r-1} and $-2,-5,\dots,\frac{-3r+2}{2}$ to v_2,v_4,\dots,v_r in that order. Consequently, designate the labels $3,6,\dots,\frac{3r-6}{2}$ corresponding to $\omega_1,\omega_3,\dots,\omega_{r-3}$ and $-3,-6,\dots,\frac{-3r+6}{2}$ to $\omega_2,\omega_4,\dots,\omega_{r-2}$ in that order. Fix the label $\frac{3r-2}{2}$ with ω_r . In each instance, the edges $\mu_iv_i,v_i\omega_i, 1 \le i \le r-1$ are are designated with 1 and the remaining edges that are unlabeled with 1. Eventually, $\overline{\mathbb{S}}_{\varphi_1} = 2r-2$ and $\overline{\mathbb{S}}_{\varphi_1^c} = 2r-1$.

Theorem 3.6. The double quadrilateral snake DQ_r is a PMC-graph for all $r \ge 2$.

Proof. Let $V(DQ_r) = \{\mu_i : 1 \le i \le r \} \cup \{\nu_i, \zeta_i, \eta_i, \omega_i \mid 1 \le i \le r - 1\}.$

Additionally $E(DQ_r) = \{u_i u_{i+1}, \mu_i v_i, v_i \zeta_i, u_{i+1} \zeta_j, u_{i+1} \omega_i, u_i \eta_i, \eta_i \omega_i : 1 \le i \le r-1\}$. Eventually, the double quadrilateral snake DQ_r consists of 7r-7 edges and 5r-4 edges.

Case (A): For odd r

Allocate the labels $-1, -6, ..., \frac{-5r+13}{2}$ corresponding to $\mu_1, \mu_3, ..., \mu_{r-2}$ and $-3, -8, ..., \frac{-5r+9}{2}$ to $\mu_2, \mu_4, ..., \mu_{r-1}$ in that order. Put the label 1 with μ_r . Subsequently, designate the labels $3, 8, ..., \frac{5r-9}{2}$ according to $\nu_1, \nu_3, ..., \nu_{r-2}$ and





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 $5,10,...,\frac{5r-5}{2}$ to $\nu_2,\nu_4,...,\nu_{r-1}$ in that order. Allocate the labels $-2,-7,...,\frac{-5r+11}{2}$ corresponding to $\zeta_1,\zeta_3,...,\zeta_{r-2}$ and $-4,-9,...,\frac{-5r+7}{2}$ to $\zeta_2,\zeta_4,...,\zeta_{r-1}$ in that order. Designate the labels $2,7,...,\frac{5r-11}{2}$ according to $\eta_1,\eta_3,...,\eta_{r-2}$ and $-5,-10,...,\frac{-5r+5}{2}$ to $\eta_2,\eta_4,...,\eta_{r-1}$ in that order. Eventually, allocate the labels $4,9,...,\frac{5r-9}{2}$ corresponding to $\omega_1, \omega_3, ..., \omega_{r-2}$ and 6,11, ..., $\frac{5r-13}{2}$ to $\omega_2, \omega_4, ..., \omega_{r-3}$ in that order. Fix 1 with the vertex ω_{r-1} . Consequently, the edges $\mu_i \nu_i, \nu_i \omega_i, 1 \leq i \leq r-1, u_i \eta_i, \tilde{u_{i+1}} \omega_i, i=1,3,...,r-2 \text{ and } \eta_i \omega_i, i=2,4,...,r-2 \text{ are labeled with 1 and the remaining edges are labeled with integers other than 1. Eventually, <math>\overline{\mathbb{S}}_{\varphi_1} = \frac{7r-7}{2} = \overline{\mathbb{S}}_{\varphi_1^c}$.

Case (B): For even r

Designate the labels $-1, -6, \dots, \frac{-5r+8}{2}$ corresponding to $\mu_1, \mu_3, \dots, \mu_{r-1}$ and $-3, -8, \dots, \frac{-5r+4}{2}$ to $\mu_2, \mu_4, \dots, \mu_r$ in that order. Consequently, allocate the labels $3, 8, \dots, \frac{5r-4}{2}$ according to v_1, v_3, \dots, v_{r-1} and $5, 10, \dots, \frac{5r-10}{2}$ to v_2, v_4, \dots, v_{r-2} in that order. Allocate the labels $-2, -7, \dots, \frac{-5r+6}{2}$ corresponding to $\zeta_1, \zeta_3, \dots, \zeta_{r-1}$ and $-4, -9, \dots, \frac{5r-12}{2}$ to $\zeta_2, \zeta_4, \dots, \zeta_{r-2}$ in that order. Designate the labels $2, 7, \dots, \frac{5r-6}{2}$ according to $\eta_1, \eta_3, \dots, \eta_{r-1}$ and $-5, -10, \dots, \frac{-5r+10}{2}$ to $\eta_2, \eta_4, \dots, \eta_r$ in that order. Eventually, allocate the labels $4, 9, \dots, \frac{5r-12}{2}$ corresponding to $\omega_1, \omega_3, \dots, \omega_{r-3}$ and $6, 11, \dots, \frac{5r-8}{2}$ to $\omega_2, \omega_4, \dots, \omega_{r-2}$ in that order. Fix 1 with the vertex ω_{r-1} . Consequently, the edges $\mu_i \nu_i, \nu_i \omega_i$, $1 \le i \le r-1$, $u_i\eta_i, 1 \leq i \leq r-1$ and i is odd, $u_{i+1}\omega_i, 1 \leq i \leq r-1$ and i is odd, and $\eta_i\omega_i, 1 \leq i \leq r-2$ and i is even and $u_i\omega_{i-1}$ are designated with 1 and the remaining edges that are unlabeled with 1. Eventually, $\overline{\mathbb{S}}_{\varphi_1} = \frac{7r-8}{2}$ and $\overline{\mathbb{S}}_{\varphi_1^c} = \frac{7r-6}{2}$.

Theorem 3.7. The alternate double quadrilateral snake $A(DQ_r)$ is a PMC-graph for all $r \ge 3$.

Case (A): Both the edges $\mu_{r-1}\mu_r$ and $\mu_1\mu_2$ are located on the double quadrilateral for even $r \ge 4$. Then $V(A(DQ_r)) = \{ \mu_i \mid 1 \le i \le r \} \cup \{ \nu_i, \zeta_i, \eta_i, \omega_i \mid 1 \le i \le \frac{r}{2} \}.$

Additionally $E(A(DQ_r)) = \{u_i u_{i+1}\} \cup \{\mu_{2i-1} v_i, v_i \zeta_i, u_{2i} \zeta_j, u_{2i} \omega_i, u_{2i-1} \eta_i, \eta_i \omega_i : 1 \le i \le \frac{r}{2}\}$. Presently, $A(DQ_r)$ includes 4r-1 edges and 3r vertices. Designate the labels $-1, -4, ..., \frac{-3r+4}{2}$ corresponding to $\mu_1, \mu_2, ..., \mu_{r-1}$ and $-3, -6, ..., \frac{-3r+2}{2}$ to $\mu_2, \mu_4, ..., \mu_r$ in that order. Consequently, allocate the labels $3, 6, ..., \frac{3r}{2}$ according to $v_1, v_2, ..., v_{r/2}$ and $-2, -5, ..., \frac{-3r+2}{2}$ to $\zeta_1, \zeta_2, ..., \zeta_{r/2}$ in that order. Designate the labels $2, 5, ..., \frac{3r-2}{2}$ according to $\eta_1, \eta_2, ..., \eta_{r/2}$ and $4, 7, ..., \frac{3r-4}{2}$ to $\omega_1, \omega_2, ..., \omega_{r-2/2}$ in that order. Fix 1 with the vertex $\omega_{r/2}$. Consequently, the edges $\mu_{2i-1}v_i, v_i\zeta_i, \mu_{2i-1}\eta_i, 1 \le i \le \frac{r}{2}$, $u_{2i}\omega_i$, $1 \le i \le \frac{r-2}{2}$ are are designated with 1 and the remaining edges that are unlabeled with 1. Eventually, $\overline{\mathbb{S}}_{\varphi_1} = 0$ 2r-1 and $\overline{\mathbb{S}}_{\varphi_1^c}=2r$.

Case (B): Both the edges $\mu_{r-1}\mu_r$ and $\mu_1\mu_2$ are not located on the double quadrilateral for odd $r \ge 4$.

Then $V(A(DQ_r)) = \{\mu_i, \nu_j, \zeta_j, \eta_j, \omega_j : 1 \le i \le r \text{ and } 1 \le j \le \frac{r-2}{2}\}$. Additionally $E(A(DQ_r)) = \{u_i u_{i+1} : 1 \le r \le r\} \cup \{u_i u_{i+1} : 1 \le r \le r\}$ $\{\mu_{2i}v_i, v_i\zeta_i, u_{2i+1}\zeta_j, u_{2i+1}\omega_i, u_{2i}\eta_i, \eta_i\omega_i: 1 \leq i \leq \frac{r-2}{2}\}. \text{ Presently, it includes } 4r-7 \text{ edges and } 3r-4 \text{ vertices. Fix 1 with } 1 \leq i \leq \frac{r-2}{2}\}.$ the vertex μ_1 . Designate the labels $-1, -4, ..., \frac{-3r+4}{2}$ corresponding to $\mu_2, \mu_4, ..., \mu_r$ and $-3, -6, ..., \frac{-3r+6}{2}$ to $\mu_3, \mu_5, ..., \mu_{r-1}$ in that order. Consequently, allocate the labels $3, 6, ..., \frac{3r-6}{2}$ according to $\nu_1, \nu_2, ..., \nu_{r-2/2}$ and $-2, -5, \dots, \frac{-3r+8}{2}$ to $\zeta_1, \zeta_2, \dots, \zeta_{r-2/2}$ in that order. Designate the labels 2,5, ..., $\frac{3r-8}{2}$ according to $\eta_1, \eta_2, \dots, \eta_{r-2/2}$ and $4,7,\ldots,\frac{3r-4}{2}$ to $\omega_1,\omega_2,\ldots,\omega_{r-2/2}$ in that order. Consequently, the edges $\mu_{2i}v_i,v_i\zeta_i,u_{2i}\eta_i,u_{2i+1}\omega_i,1\leq i\leq \frac{r-2}{2}$ are designated with 1 and the remaining edges that are unlabeled with 1. Eventually, $\overline{\mathbb{S}}_{\varphi_1} = 2r - 4$ and $\overline{\mathbb{S}}_{\varphi_1^c} = 2r - 3$.

Case (C): For odd $r \ge 3$, $\mu_{r-1}\mu_r$ and $\mu_1\mu_2$ are the edges that lie on and off the double quadrilateral, respectively. Let $(A(DQ_r)) = \{\mu_i, \nu_j, \zeta_j, \eta_j, \omega_j : 1 \le i \le r \text{ and } 1 \le j \le \frac{r-2}{2}\}.$

Additionally $E(A(DQ_r)) = \{u_i u_{i+1}: 1 \le r \le r\} \cup \{\mu_{2i} v_i, v_i \zeta_i, u_{2i+1} \zeta_j, u_{2i+1} \omega_i, u_{2i} \eta_i, \eta_i \omega_i: 1 \le i \le \frac{r-1}{2}\}.$ Presently, it includes 4r-4 edges and 3r-2 vertices. Fix 1 with the vertex μ_1 . Designate the labels $-1,-4,...,\frac{-3r+7}{2}$





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corresponding to $\mu_2, \mu_4, \dots, \mu_{r-1}$ and $-3, -6, \dots, \frac{-3r+3}{2}$ to $\mu_3, \mu_5, \dots, \mu_r$ in that order. Consequently, allocate the labels $3,6,\dots,\frac{3r-3}{2}$ according to $v_1,v_2,\dots,v_{r-1/2}$ and $-2,-5,\dots,\frac{-3r+5}{2}$ to $\zeta_1,\zeta_2,\dots,\zeta_{r-1/2}$ in that order. Designate the labels $2,5,\dots,\frac{3r-5}{2}$ according to $\eta_1,\eta_2,\dots,\eta_{r-1/2}$ and $4,7,\dots,\frac{3r-7}{2}$ to $\omega_1,\omega_2,\dots,\omega_{r-3/2}$ in that order. Fix $\frac{-3r+7}{2}$ with the vertex $\omega_{r-1/2}$. Consequently, the edges $\mu_{2i}v_i,v_i\zeta_i,\ u_{2i}\eta_i,1\leq i\leq \frac{r-1}{2},\ u_{2i+1}\omega_i,1\leq i\leq \frac{r-3}{2}$ are designated with 1 and the remaining edges that are unlabeled with 1. Eventually, $\overline{\mathbb{S}}_{\varphi_1}=2r-2$ and $\overline{\mathbb{S}}_{\varphi_1^c}=2r-2$.

Case (D): For odd $r \ge 3$, $\mu_{r-1}\mu_r$ and $\mu_1\mu_2$ are the edges that lie off and on the double quadrilateral, respectively. Let $V(A(DQ_r)) = \{\mu_i, \nu_j, \zeta_j, \eta_j, \omega_j : 1 \le i \le r \text{ and } 1 \le j \le \frac{r-2}{2}\}$

Additionally $E(A(DQ_r)) = \{u_iu_{i+1}: 1 \le r \le r\} \cup \{\mu_{2i-1}v_i, v_i\zeta_i, u_{2i}\zeta_j, u_{2i}\omega_i, u_{2i-1}\eta_i, \eta_i\omega_i: 1 \le i \le \frac{r-1}{2}\}$. Presently, it includes 4r-4 edges and 3r-2 vertices. Fix 1 with the vertex μ_r . Designate the labels $-1, -4, \dots, \frac{-3r+7}{2}$ corresponding to $\mu_1, \mu_3, \dots, \mu_{r-2}$ and $-3, -6, \dots, \frac{-3r+3}{2}$ to $\mu_3, \mu_5, \dots, \mu_{r-1}$ in that order. Consequently, label $v_i, \zeta_i, \eta_i, \omega_i$, just as in case (C). The edges $\mu_{2i-1}v_i, v_i\zeta_i, u_{2i-1}\eta_i, 1 \le i \le \frac{r-1}{2}, u_{2i}\omega_i, 1 \le i \le \frac{r-3}{2}$ and $\eta_{r-1/2}, \omega_{r-1/2}$ are designated with 1 and the remaining edges that are unlabeled with 1.Eventually, $\overline{\mathbb{S}}_{\varphi_1} = 2r-2$ and $\overline{\mathbb{S}}_{\varphi_1^c} = 2r-2$.

Theorem 3.8. The triple triangular snake TT_r is not PMC-graph for all $r \ge 2$ (except r = 3).

Proof. Let $V(TT_r) = \{\mu_i, \nu_j, \eta_j, \omega_j \mid 1 \le i \le r \text{ and } 1 \le j \le r - 1 \}.$

Additionally $E(TT_r) = \{\mu_i \nu_i, \mu_i \eta_i, \mu_i \omega_i, \mu_{i+1} \nu_i, \mu_{i+1} \eta_i, \mu_{i+1} \omega_i, \mu_i \mu_{i+1} : 1 \le i \le r-1\}$. Certainly, the triple triangular snake TT_r consists of 7r-7 edges and 4r-3 vertices.

Case (A): r = 2

Suppose the triple triangular snake TT_2 is PMC-graph. Accordingly, two possible outcomes exist if 1 is given to $\mu\nu$: both $\varphi(\mu) + \varphi(\nu) = 1$ and $\varphi(\mu) + \varphi(\nu) = 2$ can be utilized. As a result, there can be a maximum of 2 edges labeled with 1. That's $\overline{\mathbb{S}}_{\varphi_1} \leq 2$. Subsequently, $\overline{\mathbb{S}}_{\varphi_1^c} \geq 5$. Eventually $\overline{\mathbb{S}}_{\varphi_1^c} - \overline{\mathbb{S}}_{\varphi_1} \geq 5 - 2 = 3 > 1$, which contradicts itself.

Case (B): r = 3

Allocate the labels -1, -2, -3 corresponding to μ_1, μ_2, μ_3 and 1,2,3 to ν_1, η_1, ω_1 in that order. Eventually, designate the labels 4, -4,4 according to ν_2, η_2, ω_2 . Consequently, $\overline{\mathbb{S}}_{\varphi_1} = 7 = \overline{\mathbb{S}}_{\varphi_1^c}$.

Case (C): $r \ge 4$

Suppose the triple triangular snake TT_r is PMC-graph. Accordingly, two possible outcomes exist if 1 is given to $\mu\nu$: both $\varphi(\mu)+\varphi(\nu)=1$ and $\varphi(\mu)+\varphi(\nu)=2$ can be utilized. As a result, there can be a maximum of 2r+1 edges labeled with 1. That's $\overline{\mathbb{S}}_{\varphi_1} \leq 2r+1$. Subsequently, $\overline{\mathbb{S}}_{\varphi_1^c} \geq 7r-7-(2r+1)=5r-8$. Eventually $\overline{\mathbb{S}}_{\varphi_1^c}-\overline{\mathbb{S}}_{\varphi_1} \geq 5r-8-(2r+1)=3r-9=3>1$, which contradicts itself.

Theorem 3.9. The triple quadrilateral snake TQ_r is a PMC-graph for all $r \ge 2$ (except r = 3).

Proof. Let $V(TQ_r) = \{\mu_i | 1 \le i \le r \} \cup \{\nu_i, \eta_i, \omega_i, \zeta_i, \xi_i, \delta_i | 1 \le i \le r - 1\}.$

Additionally $E(TQ_r) = \{\mu_i \mu_{i+1}, \mu_i \nu_i, \mu_i \xi_i, \mu_i \omega_i, \mu_{i+1} \zeta_i, \mu_{i+1} \delta_i, \mu_{i+1} \eta_i, \nu_i \eta_i, \xi_i \delta_i, \omega_i \zeta_i: 1 \le i \le r-1 \}$. Certainly, the triple quadrilateral snake TQ_r consists of 10r-10 edges and 7r-6 vertices.

Case (A): For even r

Fix the label 3 with μ_1 . Next allocate the labels 9,16, ..., $\frac{7r-10}{2}$ corresponding to μ_3 , μ_5 , ..., μ_{r-1} and -4, -11, ..., $\frac{-7r+20}{2}$ to μ_2 , μ_4 , ..., μ_{r-2} in that order. Fix the label $\frac{-7r+8}{2}$ with μ_r . Designate the labels -1, -8, ..., $\frac{-7r+12}{2}$ according to ν_1 , ν_3 , ..., ν_{r-1} and 6,13, ..., $\frac{7r-16}{2}$ to ν_2 , ν_4 , ..., ν_{r-2} in that order. Fix 2 with the vertex η_1 . Consequently, allocate the labels 10,17, ..., $\frac{7r-8}{2}$ corresponding to η_1 , η_3 , ..., η_{r-1} and 10, 10, ..., 10, 10, ...,





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Eventually, allocate the labels 4,11, ..., $\frac{7r-6}{2}$ corresponding to $\zeta_1, \zeta_3, ..., \zeta_{r-1}$ and $-6, -13, ..., \frac{-7r+16}{2}$ to $\zeta_2, \zeta_4, ..., \zeta_{r-2}$ in that order. Designate the labels $-3, -10, ..., \frac{-7r+22}{2}$ according to $\xi_1, \xi_3, ..., \xi_{r-3}$ and $8, 15, ..., \frac{7r-12}{2}$ to $\xi_2, \xi_4, ..., \xi_{r-2}$ in that order. Allocate the labels $5, 12, ..., \frac{7r-18}{2}$ corresponding to $\delta_1, \delta_3, ..., \delta_{r-3}$ and $-7, -14, ..., \frac{-7r+14}{2}$ to $\delta_2, \delta_4, ..., \delta_{r-2}$ in that order. Fix the label $\frac{-7r+6}{2}$ with δ_{r-1} . Subsequently, the edges $\mu_i \nu_i, \nu_i \eta_i, \xi_i \delta_i$, for $1 \le i \le r-1$, $\omega_i \zeta_i, \mu_{i+1} \zeta_j$ for $1 \le i \le r-2$, $\mu_r \delta_{r-1}$ and $\mu_1 \xi_1$ are designated with 1 and the remaining edges that are unlabeled with 1.

Case (B): For odd r

Fix the label 3 with μ_1 . Next allocate the labels 9,16, ..., $\frac{7r-17}{2}$ corresponding to μ_3 , μ_5 , ..., μ_{r-2} and -4, -11, ..., $\frac{-7r+13}{2}$ to μ_2 , μ_4 , ..., μ_{r-1} in that order. Fix the label $\frac{7r-11}{2}$ with μ_r . Designate the labels -1, -8, ..., $\frac{-7r+19}{2}$ according to ν_1 , ν_3 , ..., ν_{r-2} and 6,13, ..., $\frac{7r-9}{2}$ to ν_2 , ν_4 , ..., ν_{r-1} in that order. Fix 2 with the vertex η_1 . Consequently, allocate the labels 10,17, ..., $\frac{7r-15}{2}$ corresponding to η_1 , η_3 , ..., η_{r-2} and -5, -12, ..., $\frac{-7r+11}{2}$ to η_2 , η_4 , ..., η_{r-1} in that order. Designate the labels -2, -9, ..., $\frac{-7r+17}{2}$ according to ω_1 , ω_3 , ..., ω_{r-2} and 7,14, ..., $\frac{7r-19}{2}$ to ω_2 , ω_4 , ..., ω_{r-1} in that order. Eventually, allocate the labels 4,11, ..., $\frac{7r-13}{2}$ corresponding to ζ_1 , ζ_3 , ..., ζ_{r-2} and -6, -13, ..., $\frac{-7r+9}{2}$ to ζ_2 , ζ_4 , ..., ζ_{r-1} in that order. Designate the labels -3, -10, ..., $\frac{-7r+15}{2}$ according to ξ_1 , ξ_3 , ..., ξ_{r-2} and 8,15, ..., $\frac{-7r+9}{2}$ to ξ_2 , ξ_4 , ..., ξ_{r-3} in that order. Fix the label 1 with ξ_{r-1} . Allocate the labels 5,12, ..., $\frac{7r-11}{2}$ corresponding to δ_1 , δ_3 , ..., δ_{r-2} and -7, -14, ..., $\frac{-7r+7}{2}$ to δ_2 , δ_4 , ..., δ_{r-1} in that order. Subsequently, the edges $\mu_i \nu_i$, $\nu_i \eta_i$, $\xi_i \delta_i$, for $1 \le i \le r-1$, $\omega_i \zeta_i$, $\mu_{i+1} \zeta_j$ for $1 \le i \le r-2$, $\mu_{r-1} \mu_r$ and $\mu_1 \xi_1$ are designated with 1 and the remaining edges that are unlabeled with 1. In each instance, $\overline{\mathbb{S}}_{\varphi_1} = 5r-5 = \overline{\mathbb{S}}_{\varphi_1}^c$.

CONCLUSION

In this article, we have examined the PMC- labeling behavior of some snake graphs. There is unfinished business in terms of studying related theorems for other graphs.

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RESEARCH ARTICLE

Some Properties of b# Interior and b#-Closure in Intuitionistic **Topological Spaces**

C. Velraj¹ and R. Jamuna Rani²

¹Research Scholar, Register No. 20221172091005, Department of Mathematics, Rani Anna Govt. College for Women, Tirunelveli. Affiliated to MS University, Tirunelveli

²Asst. Professor and Head of the Dept. of Mathematics, Govt. Arts and Science College (Women), Sathankulam.

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*Address for Correspondence

C. Velraj

Research Scholar, Department of Mathematics, Rani Anna Govt. College for Women, Tirunelveli. Affiliated to MS University, Tirunelveli

Email: velraj.c79@gmail.com



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ABSTRACT

This paper's main goal is to present intuitionistic topological spaces, specifically b#-closure and b#interior. Numerous attributes and traits are listed.

Keywords: Intuitionistic b*interior, Intuitionistic b*closure.

INTRODUCTION AND PRELIMINARIES

Numerous generalizations of this core idea have been made since Zadeh[1] introduced fuzzy sets. The idea of "intuitionistic sets" presented by Coker[2] in 1996. All of the sets in this discrete version of the intuitionistic fuzzy set are completely crisp sets. He has investigated a few basic topological characteristics on intuitionistic sets. Because of their importance to general topology, open sets are currently a hot topic for study among academics all over the world. In intuitionistic topological spaces, semi open sets, pre open sets, and b-open sets were researched by Sasikala[4,5,6] and Navaneetha Krishnan. The b* open sets are researched by Velraj [7]. In topological spaces, Gnanambal Ilango[8] possesses the qualities of α -interior and α -closure. This work introduces and characterizes the features of intuitionistic b* open sets. Numerous pertinent papers have already been published in books. In this study, we described some of the features of b# interior and b# closure in intuitionistic topological spaces.





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Definition1.1*cited from* [3]: A family τ of IS's in \widetilde{Y} satisfyingthefollowingaxioms is called an intuitionistic topology.

- (T₁) $\widetilde{\phi}$, $\widetilde{Y} \in \tau$,
- (T₂) $\tilde{G}_1 \cap \tilde{G}_2 \in \tau$ for any \tilde{G}_1 , $\tilde{G}_2 \in \tau$ and
- (T₃) $\cup \tilde{G}_i \in \tau$ for any arbitrary family $\{\tilde{G}_i: i \in J\} \subseteq \tau$.

Any IS in τ is referred to as an intuitionistic open set (IOS) in \widetilde{Y} and the pair (\widetilde{Y} , τ) is called an intuitionistic topological space (ITS).

Definition 1.2*cited from*[3,9]: A subset \tilde{S} in (\tilde{Y},τ) is called

- i) Intuitionistic regular open [3](IRO)*if* $\widetilde{S} = int(cl(\widetilde{S}))$
- ii) Intuitionistic b# open [9](Ib#O) $if\tilde{S} = cl(int(\tilde{S})) \bigcup int(cl(\tilde{S}))$

Lemma 1.3*cited from* [10]:For a subset \tilde{S} of an ITS (\tilde{Y}, τ) ,

- (i) $int(Ibcl(\tilde{S}))=Ibcl(int(\tilde{S}))=int(cl(int(\tilde{S}))).$
- (ii) $cl(\operatorname{Ibint}(\widetilde{S}))=\operatorname{Ibint}(cl(\widetilde{S}))=cl(\operatorname{int}(cl(\widetilde{S}))).$
- (iii) $Ibint(\tilde{S}) = Isint(\tilde{S}) \bigcup Ipint(\tilde{S}).$

Intuitionistic b[#] interior

We define intuitionistic b# interior and go over its characteristics in this section.

Definition 2.1. Consider the ITS(\widetilde{Y} , τ). The union of all intuitionistic b^* -open sets contained in \widetilde{U} is called the intuitionistic b^* interior of an intuitionistic set \widetilde{U} and is denoted by $Ib^*int(\widetilde{U})$.

That is $\operatorname{Ib}^{\sharp}int(\widetilde{U}) = \bigcup \{\widetilde{V} \colon \widetilde{V} \subseteq \widetilde{U} \text{ and } \widetilde{V} \text{ is } \operatorname{Ib}^{\sharp} \text{ open} \}.$

The qualities of intuitionistic b* interior are provided in the following Propositions.

Proposition 2.2.

- (i) $\operatorname{Ib}^{\sharp} \operatorname{int}(\tilde{\phi}) = \tilde{\phi}.$
- (ii) $\operatorname{Ib}^{\sharp} \operatorname{in} t(\widetilde{Y}) = \widetilde{Y}.$
- (iii) $\operatorname{Ib}^{\sharp}int(\widetilde{U}) \subseteq \widetilde{U}$.
- (iv) If \widetilde{U} is $Ib^{\#}$ open then $Ib^{\#}int(\widetilde{U}) = \widetilde{U}$.

Proof: The definition provides proof.

Proposition 2.3. Let (\widetilde{Y}, τ) be an ITS. Then for any two sub sets \widetilde{C} and \widetilde{D} of \widetilde{Y} , we get

- (i) If $\tilde{C} \subseteq \tilde{D}$ then $lb^{\#}int(\tilde{C}) \subseteq lb^{\#}int(\tilde{D})$.
- (ii) $\operatorname{Ib}^{\sharp}int(\operatorname{Ib}^{\sharp}int(\widetilde{C}))=\operatorname{Ib}^{\sharp}int(\widetilde{D}).$
- (iii) $\operatorname{Ib}^{\sharp}int(\widetilde{C} \bigcup \widetilde{D}) \subseteq \operatorname{Ib}^{\sharp}int(\widetilde{C}) \cap \operatorname{Ib}^{\sharp}int(\widetilde{D}).$
- (iv) $\operatorname{Ib}^{\sharp}int(\widetilde{C} \bigcup \widetilde{D}) \supset \operatorname{Ib}^{\sharp}int(\widetilde{C}) \bigcup \operatorname{Ib}^{\sharp}int(\widetilde{D}).$
- (v) $\operatorname{Ib}^{\sharp}int(\tilde{C}) \subset \operatorname{Ib}int(\tilde{C}).$
- (vi) $\operatorname{Ib}^{\sharp}int(\tilde{C}) \subseteq \operatorname{Isint}(\tilde{C}) \bigcup \operatorname{Ipint}(\tilde{C}).$

 $\boldsymbol{Proof.}$ Proof of (i), (iii), (iv) and (v) follows from the Definition.

Now to prove (ii). Using Proposition 2.2(iii), $\mathsf{Ib}^*int(\tilde{C}) \subseteq \tilde{C}$. Again Using (i), $\mathsf{Ib}^*int(\mathsf{b}^*int(\tilde{C})) \subseteq \mathsf{Ib}^*int(\tilde{C})$. On the other hand let $x \in \mathsf{Ib}^*int(\tilde{C})$. This implies $x \in \widetilde{D} \subseteq \tilde{C}$ for some Ib^* -open set \widetilde{D} . Now since \widetilde{D} is Ib^* -open, Ib^* -int(\widetilde{D})= \widetilde{D} . Since $\widetilde{D} \subseteq \tilde{C}$, by (i), $\mathsf{Ib}^*int(\widetilde{D}) \subseteq \mathsf{Ib}^*int(\tilde{C})$ that is $\widetilde{D} \subseteq \mathsf{Ib}^*int(\tilde{C})$ that implies $x \in \widetilde{D} \subseteq \mathsf{Ib}^*int(\tilde{C})$. This gives $x \in \mathsf{Ib}^*int(\mathsf{Ib}^*int(\tilde{C}))$. Therefore $\mathsf{Ib}^*int(\widetilde{D}) \subseteq \mathsf{Ib}^*int(\mathsf{Ib}^*int(\tilde{C}))$. This proves (ii).

Now to prove (vi). By using Lemma 1.3(iii), $Ibint(\tilde{C}) = Isint(\tilde{C}) \bigcup Ipint(\tilde{C})$. This together with (v) implies, $Ib^*int(\tilde{C}) = Isint(\tilde{C}) \cup Ipint(\tilde{C})$. This proves (vi).

It is possible to create examples to demonstrate that

(i) $\operatorname{Ib}^{\sharp}int(\widetilde{U}) \subseteq \operatorname{Ib}^{\sharp}int(\widetilde{V})$ does not imply $\widetilde{U} \subseteq \widetilde{V}$.





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- (ii) $\operatorname{Ib}^{\sharp}int(\widetilde{U} \cap \widetilde{V}) \neq \operatorname{Ib}^{\sharp}int(\widetilde{U}) \cap \operatorname{Ib}^{\sharp}int(\widetilde{V}).$
- (iii) $\operatorname{Ib}^{\sharp}int(\widetilde{U}\bigcup\widetilde{V})\neq\operatorname{Ib}^{\sharp}int(\widetilde{U})\bigcup\operatorname{Ib}^{\sharp}int(\widetilde{V}).$
- (iv) $\operatorname{Ib}^{\sharp}int(\widetilde{U}) \neq \operatorname{Ib}int(\widetilde{U}).$
- (v) $\operatorname{Ib}^{\sharp}int(\widetilde{U}) \neq \operatorname{Isint}(\widetilde{U}) \bigcup \operatorname{Ipint}(\widetilde{U}).$

Example 2.4: Let $\widetilde{Y} = \{4,3,2,1\}$ and $\tau = \{\widetilde{\phi},\widetilde{Y_1},\widetilde{Y_2},\widetilde{Y_3},\widetilde{Y}\}$ where $\widetilde{Y_1} = \langle Y, \{2,1\},\{4,3\}\rangle$, $\widetilde{Y_2} = \langle Y,\{3\}, \{4,2,1\}\rangle$, $\widetilde{a} = \langle Y,\{3,2,1\},\{4\}\rangle$. Let $\widetilde{a} = \langle Y,\{4,3\},\{1\}\rangle$ and $\widetilde{a} = \langle Y,\{4,3,1\},\{2\}\rangle$ be the intuitionistic subsets of $\widetilde{A}(\tau)$. Now $Ib^{\#}int(\widetilde{U}) = \langle Y,\{4,3\},\{2,1\}\rangle$ and $Ib^{\#}int(\widetilde{V}) = \langle Y,\{4,3\},\{2,1\}\rangle$. This shows that $Ib^{\#}int(\widetilde{U}) \subseteq Ib^{\#}int(\widetilde{V})$ does not imply $\widetilde{U} \subseteq \widetilde{V}$.

Example 2.5: Let $\widetilde{Y} = \{3,2,1\}$ and $\tau = \{\widetilde{\phi}, \widetilde{Y}_1, \widetilde{Y}_2, \widetilde{Y}_3, \widetilde{Y}\}$ where $\widetilde{Y}_1 = \langle Y, \{1\}, \{3,2\} \rangle$, $\widetilde{Y}_2 = \langle Y, \{3\}, \{2,1\} \rangle$, $\widetilde{Y}_3 = \langle Y, \{3,1\}, \{2\} \rangle$. Let $\widetilde{C} = \langle Y, \{2,1\}, \{\phi\} \rangle$ and $\widetilde{D} = \langle Y, \{3,2\}, \{\phi\} \rangle$ be the intuitionistic subsets of $(\widetilde{Y}, \tau).\widetilde{C} \cap \widetilde{D} = \langle X, \{2\}, \{\phi\} \rangle$ $Ib\#int(\widetilde{C} \cap \widetilde{D}) = \phi.$ Now, $Ib\#int(\widetilde{C}) = \langle Y, \{2,1\}, \{3\} \rangle$ and $Ib\#int(\widetilde{D}) = \langle Y, \{3,2\}, \{1\} \rangle.$ $Ib\#int(\widetilde{C}) \cap Ib\#int(\widetilde{D}) = \langle Y, \{2,1\}, \{3\} \rangle.$ Therefore, $Ib\#int(\widetilde{C} \cap \widetilde{D}) \neq Ib\#int(\widetilde{C}) \cap Ib\#int(\widetilde{D})$.

Example 2.6: Let $\widetilde{Y} = \{4,3,2,1\}$ and $\tau = \{\widetilde{\phi},\widetilde{Y_1},\widetilde{Y_2},\widetilde{Y_3},\widetilde{Y_4},\widetilde{Y_5},\widetilde{Y_6},\widetilde{Y_7},\widetilde{Y_8},\widetilde{Y}\}$ where $\widetilde{Y_1} = \langle Y, \{3,2,1\}, \{\phi\} \rangle$, $\widetilde{Y_2} = \langle Y, \{3,2\}, \{4\} \rangle$, $\widetilde{Y_3} = \langle Y, \{2,1\}, \{3\} \rangle$, $\widetilde{Y_4} = \langle Y, \{2\}, \{4,3\} \rangle$, $\widetilde{Y_5} = \langle Y, \{3\}, \{4,1\} \rangle$, $\widetilde{Y_6} = \langle Y, \{\phi\}, \{4,3,1\} \rangle$, $\widetilde{Y_7} = \langle Y, \{1\}, \{3\} \rangle$, $\widetilde{Y_8} = \langle Y, \{\phi\}, \{4,3\} \rangle$. Let $\widetilde{C} = \langle Y, \{4,3\}, \{1\} \rangle$ and $\widetilde{D} = \langle Y, \{4,1\}, \{\phi\} \rangle$ be the intuitionistic subsets of (\widetilde{Y}, τ) . $\widetilde{C} \cup \widetilde{D} = \langle X, \{4,3,1\}, \{\phi\} \rangle$ $Ib^{\#}int(\widetilde{C}) = \langle X, \{3\}, \{1\} \rangle$ and $Ib^{\#}int(\widetilde{D}) = \langle Y, \{4,1\}, \{3\} \rangle$. Now, $Ib^{\#}int(\widetilde{C}) = \langle X, \{3\}, \{1\} \rangle$ and $Ib^{\#}int(\widetilde{D}) = \langle Y, \{4,3\}, \{1\}, \{\phi\} \rangle$. Therefore $Ib^{\#}int(\widetilde{C}) \cup Ib^{\#}int(\widetilde{D})$.

Example 2.7:Let us takethe ITS in Example 2.5. Let $\widetilde{U} = \langle Y, \{2,1\}, \{\phi\} \rangle$ be an intuitionistic subset of (\widetilde{Y}, τ) . Now, $Ib^*int(\widetilde{U}) = \langle Y, \{2,1\}, \{3\} \rangle$ and $Ibint(\widetilde{U}) = \langle Y, \{2,1\}, \{\phi\} \rangle$. This shows that $Ib^*int(\widetilde{U}) \neq Ibint(\widetilde{U})$

Example 2.8:Let us take the ITS in Example 2.4. Let $\widetilde{U} = \langle Y, \{4,3\}, \{2\} \rangle$ be the intuitionistic subset of $(\widetilde{Y}, \tau).lb^*int(\widetilde{U}) = \langle Y, \{4,3\}, \{2,1\} \rangle$.

Now $Isint(\tilde{U}) = \langle Y, \{4,3\}, \{2\} \rangle$ and $Ipint(\tilde{U}) = \langle Y, \{3\}, \{2\} \rangle$.

Now $Isint(\widetilde{U}) \cup Ipint(\widetilde{U}) = \langle Y, \{4,3\}, \{2\} \rangle$.

This shows that $Ib^{\#}int(\widetilde{U}) \neq Isint(U) \cup Ipint(\widetilde{U})$.

Intuitionistic b*-Closure

This section presents the idea of intuitionistic b*-closure and discusses its characteristics.

Definition 3.1. Let (\tilde{Y}, τ) *be an ITS*. The intersection of all intuitionistic b^* -closed sets containing \tilde{C} is called the intuitionistic b^* -interior of an intuitionistic set \tilde{C} *in* (Y, τ) , and is denoted by $Ib^*cl(\tilde{C})$.

That is $\mathrm{Ib}^{\sharp}cl(\widetilde{C}) = \bigcap \{\widetilde{D} : \widetilde{C} \subseteq \widetilde{D} \text{ and } \widetilde{D} \text{ is } \mathrm{Ib}^{\sharp}\text{-closed}\}.$

Proposition 3.2. Let (\widetilde{Y}, τ) be an ITS. Then for any subset \widetilde{C} of \widetilde{Y} we have

- (i) $\widetilde{Y} \setminus Ib^{\#}int(\widetilde{C}) = Ib^{\#}cl(\widetilde{Y} \setminus \widetilde{C}).$
- (ii) $\widetilde{Y} \setminus \operatorname{Ib}^{\sharp} cl(\widetilde{C}) = \operatorname{Ib}^{\sharp} int(\widetilde{Y} \setminus \widetilde{C}).$

Proof. By using Definition 2.1, $\mathrm{Ib}^{\sharp}int(\widetilde{C}) = \bigcup \{\widetilde{D}: \widetilde{D} \subset \widetilde{C} \text{ and } \widetilde{D} \text{ is } \mathrm{Ib}^{\sharp}\text{-open}\}.$

Accepting reciprocity on both ends, we obtain $\widetilde{Y} \setminus Ib^*int(\widetilde{C}) = \widetilde{Y} \setminus [\bigcup \widetilde{D} : \widetilde{D} \subseteq \widetilde{C} \text{ and } \widetilde{D} \text{ is } Ib^*\text{-open}] = \bigcap \{\widetilde{Y} \setminus \widetilde{D} : \widetilde{Y} \setminus \widetilde{C} \subseteq \widetilde{Y} \setminus \widetilde{D} \text{ and } \widetilde{Y} \setminus \widetilde{D} \text{ is } Ib^*\text{-closed}\} = Ib^*cl(\widetilde{Y} \setminus \widetilde{C}). \text{ This proves (i).}$

Using(i), $\widetilde{Y} \setminus Ib^{\#int}(\widetilde{Y} \setminus \widetilde{C}) = Ib^{\#cl}(\widetilde{Y} \setminus (\widetilde{Y} \setminus \widetilde{C})) = Ib^{\#cl}(\widetilde{C})$. This implies that $Ib^{\#int}(\widetilde{Y} \setminus \widetilde{C}) = \widetilde{Y} \setminus Ib^{\#cl}(\widetilde{C})$.

Preposition3.3.

- (i) $\operatorname{Ib}^{\sharp} cl(\tilde{\phi}) = \tilde{\phi},$
- (ii) $\operatorname{Ib}^{\sharp} cl(\widetilde{Y}) = \widetilde{Y}.$
- (iii) $\widetilde{U} \subseteq \operatorname{Ib}^{\sharp} cl(\widetilde{U}).$
- (iv) If \widetilde{U} is Ib#closed then Ib# $cl(\widetilde{U}) = \widetilde{U}$.





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Proof. The definition provides proof.

Proposition 3.4. Let (\widetilde{Y}, τ) be an ITS. Then for any two sub sets \widetilde{C} and \widetilde{D} of \widetilde{Y} we have

- (i) If $\tilde{C} \subseteq \tilde{D}$ then $Ib^{\#}cl(\tilde{C}) \subseteq Ib^{\#}cl(\tilde{D})$.
- (ii) $\operatorname{Ib}^{\sharp} cl(\operatorname{Ib}^{\sharp} cl(\widetilde{C})) = \operatorname{Ib}^{\sharp} cl(\widetilde{C}).$
- (iii) $\operatorname{Ib}^{\sharp}cl(\widetilde{C}\bigcup\widetilde{D})\supseteq\operatorname{Ib}^{\sharp}cl(\widetilde{C})\bigcup\operatorname{Ib}^{\sharp}cl(\widetilde{D}).$
- (iv) $\operatorname{Ib}^{\sharp} cl(\widetilde{C} \cap \widetilde{D}) \subseteq \operatorname{Ib}^{\sharp} cl(\widetilde{C}) \cap \operatorname{Ib}^{\sharp} cl(\widetilde{D}).$
- (v) $\operatorname{Ib}^{\sharp} cl(\tilde{C}) \supseteq \operatorname{Ib} cl(\tilde{C})$.
- (vi) $\operatorname{Ib}^{\sharp}cl(\tilde{C}) \supseteq \operatorname{Iscl}(\tilde{C}) \cap \operatorname{Ipcl}(\tilde{C}).$

Proof. Proof of (i), (iii), (iv) and (v) are follows from the Definition.

Now to prove (ii). By using Proposition 2.3(ii), $Ib^*int(Ib^*int(\tilde{C}))=Ib^*-int(\tilde{C})$. Accepting reciprocity on both ends, we obtain $Ib^*cl(Ib^*cl(\tilde{C}))=Ib^*cl(\tilde{C})$. This proves (ii).

Now to prove (vi). Since by Lemma 1.3(iii), implies that $Ibcl(\tilde{C})=Iscl(\tilde{C})\bigcap Ipcl(\tilde{C})\subseteq Ib^*cl(\tilde{C})$. This proves (vi)

It is possible to create examples to demonstrate that

- (i) $\operatorname{Ib}^{\sharp}cl(\widetilde{U})\subseteq\operatorname{Ib}^{\sharp}cl(\widetilde{V})$ does not imply $\widetilde{U}\subseteq\widetilde{V}$.
- (ii) $\operatorname{Ib}^{\sharp}cl(\widetilde{U}\bigcup\widetilde{V})\neq\operatorname{Ib}^{\sharp}cl(\widetilde{U})\bigcup\operatorname{Ib}^{\sharp}cl(\widetilde{V}).$
- (iii) $\operatorname{Ib}^{\sharp}cl(\widetilde{U} \cap \widetilde{V}) \neq \operatorname{Ib}^{\sharp}cl(\widetilde{U}) \cap \operatorname{Ib}^{\sharp}cl(\widetilde{V}).$
- (iv) $Ib^{\sharp}cl(\widetilde{U}) \neq Ibcl(\widetilde{U}).$
- (v) $Ib^{\sharp}cl(\widetilde{U}) \neq Iscl(\widetilde{U}) \cap Ipcl(\widetilde{U}).$

Example 3.5:Let us take the ITS in Example 2.6. Let $\widetilde{U} = \langle Y, \{1\}, \{3,2\} \rangle$ and $\widetilde{V} = \langle Y, \{1\}, \{4,3\} \rangle$ be the intuitionistic subsets of (\widetilde{Y}, τ) . Now $lb^\sharp cl(\widetilde{U}) = \langle Y, \{1\}, \{3\} \rangle$ and $lb^\sharp cl(\widetilde{V}) = \langle Y, \{1\}, \{3\} \rangle$. This shows that $lb^\sharp cl(\widetilde{U}) \subseteq lb^\sharp cl(\widetilde{V})$ does not imply $\widetilde{U} \subseteq \widetilde{V}$.

Example 3.6: Let us takethe ITS in Example 2.5. Let $\tilde{C} = \langle Y, \{3\}, \{1\} \rangle$ and $\tilde{D} = \langle Y, \{1\}, \{2\} \rangle$ be the intuitionistic subsets of (\tilde{Y}, τ) . $\tilde{C} \cup \tilde{D} = \langle Y, \{3,1\}, \{\phi\} \rangle$. $Ib^*cl(\tilde{C} \cup \tilde{D}) = \tilde{Y}$. Now, $Ib^*cl(\tilde{C}) = \langle Y, \{3\}, \{2,1\} \rangle$ and $Ib^*cl(\tilde{D}) = \langle Y, \{3,1\}, \{2\} \rangle$. Therefore $Ib^*cl(\tilde{C} \cup \tilde{D}) \neq Ib^*cl(\tilde{C}) \cup Ib^*cl(\tilde{D})$.

Example 3.7:Let us take the Example 2.4 Let $\widetilde{C} = \langle Y, \{1\}, \{4,3\} \rangle$ and $\widetilde{D} = \langle Y, \{2\}, \{4,3\} \rangle$ be the intuitionistic subset of (\widetilde{Y}, τ) . $\widetilde{C} \cap \widetilde{D} = \phi$. $Ib^{\varepsilon}cl(\widetilde{C} \cap \widetilde{D}) = \phi$. Now, $Ib^{\varepsilon}cl(\widetilde{C}) = \langle Y, \{2,1\}, \{4,3\} \rangle$ and $Ib^{\varepsilon}cl(\widetilde{D}) = \langle Y, \{2,1\}, \{4,3\} \rangle$. $Ib^{\varepsilon}cl(\widetilde{C}) \cap Ib^{\varepsilon}cl(\widetilde{C}) \cap Ib^{\varepsilon}cl(\widetilde{C}) \cap Ib^{\varepsilon}cl(\widetilde{C})$.

Example 3.8:Let us take the ITS in Example 2.6. Let $\widetilde{U} = \langle Y, \{1\}, \{4,3,2\} \rangle$ be the intuitionistic subset of (\widetilde{Y}, τ) . Now $Ib^{\sharp}cl(\widetilde{U}) = \langle Y, \{1\}, \{3\} \rangle$. This shows that $Ib^{\sharp}cl(\widetilde{U}) \neq Ibcl(\widetilde{U})$.

Example 3.9:Let us take the ITS in Example 2.4. Let $\widetilde{U} = \langle Y, \{4,3\}, \{2\} \rangle$ be the intuitionistic subset of $(\widetilde{Y}, \tau).lb^*cl(\widetilde{U}) = \langle Y, \{2,1\}, \{4,3\} \rangle$.

Now $Iscl(\widetilde{U}) = \langle Y, \{2,1\}, \{4,3\} \rangle$ and $Ipcl(\widetilde{U}) = \langle Y, \{2\}, \{4,3,1\} \rangle$.

Now $Iscl(\widetilde{U}) \cap Ipcl(\widetilde{U}) = \langle Y, \{2\}, \{4,3,1\} \rangle$.

This shows that $Ib^{\sharp}cl(\widetilde{U}) \neq Iscl(\widetilde{U}) \cap Ipcl(\widetilde{U})$.

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RESEARCH ARTICLE

On the Computation of the Metric Dimension of Unit Graph

S.Nithya¹ and V.Prisci²

- ¹Assistant Professor, St. Xavier's College (Autonomous), Palayamkottai-627002, Tamil Nadu, India.
- ²Reg.No.: 20211282092020, St. Xavier's college (Autonomous), Palayamkottai-627002.
- ²Manonmanium Sundaranar University, Abishekapatti, Tirunelveli-627012, Tamil Nadu, India.

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*Address for Correspondence S.Nithya

Assistant Professor, St. Xavier's College (Autonomous), Palayamkottai-627002, Tamil Nadu, India. Email: nithyasxc@gmail.com



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ABSTRACT

To determine the unit graph's metric dimension for finite commutative rings. We explore the metric dimension of the unit graph by finding the minimal cardinality of the resolving set. We examine the unit graph's metric dimension for both the product of the ring of integers modulo α , where α is prime and the ring of integers modulo n, where n is natural numbers. Finding the bare minimum of stations and pinpointing their locations will help address the navigation challenge in space stations. Here, we explore the novel notion of metric dimension in unit graph by calculating the minimum number of resolving set.

Keywords: Metric dimension, unit graph

INTRODUCTION

The unit graph $G(Z_n)$ was first introduced by R.P.Grimaldi in 1990 [1] and it was extended to $G(\mathcal{R})$ by N.Ashrafi and others [2]. Following that, a large number of mathematicians examined various aspects of unit graphs of commutative rings in terms of graph theory. Similar concepts has been studied in [3],[4]. Slater.P.J established the idea of metric dimension initially[5]. In this paper, we discuss the metric dimension of the unit graph of integers modulo n and product of integers modulo α where α is prime.

METHODOLOGY





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In this study, we will refer to the commutative ring with identity as \mathcal{R} . Consider a ring \mathcal{R} with $U(\mathcal{R})$ as its set of unit elements. The graph $G(\mathcal{R})$ is said to be the *unit graph* with all the elements of \mathcal{R} to be the vertices if for any two distinct vertices $s, t \in \mathcal{R}$, the vertices s and t are adjacent if and only if $s + t \in U(\mathcal{R})$. For an ordered subset $\Psi =$ $\{\psi_1, \psi_2, ..., \psi_k\}$ of vertices in a connected graph G and a vertex s of G, the metric representation of with respect to Ψ is the k-vector ordered set, $\zeta(s|\Psi) = (d(s,\psi_1),d(s,\psi_2),...,d(s,\psi_k))$. The set Ψ is a resolving set for G if $\zeta(s|\Psi) =$ $\zeta(t|\Psi)$ implies that t=t for all pairs t=t for all pairs t=t for all pairs t=t for t=t for t=t for t=t for all pairs t=t for all represented by the metric dimension, $\beta(G)$.

RESULTS AND DISCUSSION

Theorem 3.1.Let \mathcal{R} be a ring. Then $\beta(G(\mathcal{R})) = |\mathcal{R}| - 1$ if and only if \mathcal{R} is a division ring with Char $\mathcal{R} = 2$.

Proof: The result follows from Theorem 3.4 of [2] and Lemma 2.2 of [6].

Theorem 3.2. Let \mathcal{R} be a ring with maximal ideal \mathfrak{M} such that $\left|\frac{R}{\mathfrak{m}}\right| = 2$ or n-1. Then $\beta(G(\mathcal{R})) = |\mathcal{R}| - 2$ if and only if \mathcal{R} is a

Proof: The result follows from Theorem 3.5 of [2] and Corollary 2.1 of [6].

Theorem 3.3. Let α be a prime number. If $n = 2\alpha$, then $\beta(G(\mathbb{Z}_n)) = 2$ or n - 1.

Proof. If $\alpha = 2$, then $\beta(G(\mathbb{Z}_4)) = 2$ since $G(\mathbb{Z}_4)$ is a cycle. If $\alpha > 2$, then $G(\mathbb{Z}_{2\alpha})$ is isomorpic to a $\alpha - 1$ regular bipartite graph. Hence by Theorem 1 of [7], the result follows.

Theorem 3.4. Let α be a prime number, then $\beta(G(\mathbb{Z}_{\alpha})) = \begin{cases} \frac{\alpha-1}{2}, & \text{if } \alpha > 2\\ 1, & \text{if } \alpha = 2 \end{cases}$ Proof. If $\alpha = 2$, $\beta(G(\mathbb{Z}_2)) = 1$, since $G(\mathbb{Z}_2)$ is a path. If $\alpha > 2$, let $V_0 = \{0\}$, $V_j = \{j, \alpha - j\}$, $1 \le j \le \frac{\alpha-1}{2}$ forms the distance similar classes. Consider $\Psi = \{j\}$, $1 \le j \le \frac{\alpha-1}{2}$. Also, $0 \notin \Psi$ since $\zeta(0|\Psi) = (1,1,...,1)$ is a unique metric representation. Hence, by Theorem 2.1. $\zeta(G(\mathbb{Z}_{\alpha})) = \frac{\alpha-1}{2}$ representation. Hence, by Theorem 2.1 of [6], $\beta(G(\mathbb{Z}_{\alpha})) = \frac{\alpha-1}{2}$.

Theorem 3.5. Let $\mathcal{R} \cong \mathbb{Z}_{\alpha} \times \mathbb{Z}_{\beta}$ where α and β are prime. If $\alpha = \beta, \alpha, \beta \geq 3$, then the metric dimension of the unit graph, $\beta(G(\mathcal{R})) \leq \frac{3}{2}(\alpha - 1).$

 $\textit{Proof.} \ \text{Let us define} \Psi_1 = \{(0,1), (1,0), (\alpha-1,\alpha-1)\}, \Psi_2 = \{(2,3), (4,5), \dots, (\alpha-3,\alpha-2)\},$

 $\Psi_3 = \{(3,2), (5,4), \dots, (\alpha-2, \alpha-3)\}, \Psi_4 = \{(3,3), (5,5), \dots, (\alpha-2, \alpha-2)\}.$ From the construction of the above sets,

$$\Psi_1, \Psi_2, \Psi_3 \text{ and } \Psi_4 \text{ are mutually disjoint to each other. Let } \Psi = \Psi_1 \cup \Psi_2 \cup \Psi_3 \cup \Psi_4.$$
 In Fig.1, for (k, l) , $(m, n) \in \mathcal{R}$, $d\big((k, l), (m, n)\big) = \begin{cases} 2, & \text{if } k + m \in U(R) \text{ or } l + n \in U(R) \\ 1, & \text{otherwise} \end{cases}$

The following metric representation are observed from Fig.1,

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\zeta((0,0)|\Psi) = (2,2,1,1,1,...,1,1,1,...,1,1,1,...,1), \zeta((0,2)|\Psi) = (2,1,1,1,1,...,2,1,1,...,1,1,1,...,2),
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$$\zeta((0,3)|\Psi) = (2,1,1,1,1,...,1,1,1,...,2,1,1,...,1),...,\zeta((0,\alpha-2)|\Psi) = (2,1,1,1,1,...,1,2,1,...,1,1,1,...,1),$$

$$\zeta((0,\alpha-1)|\Psi) = (2,1,1,1,1,\dots,1,1,1,\dots,1,1,1,\dots,1), \zeta((1,1)|\Psi) = (1,1,2,1,1,\dots,1,1,1,\dots,1,1,1,\dots,1),$$

$$\zeta((1,2)|\Psi) = 1,1,2,1,1,...,2,1,1,...,1,1,1,...2),...,\zeta((1,\alpha-2)|\Psi) = (1,1,2,1,1,...,1,2,1,...1,1,1,...1),$$

$$\zeta((1,\alpha-1)|\Psi) = (2,1,2,1,1,\dots,1,1,1,\dots,1,1,1,\dots,1), \zeta((2,0)|\Psi) = (1,2,1,1,1,\dots,1,1,1,\dots,2,1,1,\dots,2),$$

$$\zeta((2,1)|\Psi) = (1,1,2,1,1,...,1,1,1,...,2,1,1,...,2), \zeta((2,2)|\Psi) = (1,1,1,1,1,...,2,1,1,...,2,1,1,...,2),...,$$

$$\zeta((2,\alpha-1)|\Psi) = (2,1,1,1,1,...,1,1,1,...,2,1,1,...,2), \zeta((\alpha-1,0)|\Psi) = (1,2,1,1,1,...,1,1,1,...,1,1,1,...,1),$$

$$\zeta((\alpha-1,1)|\Psi)=(1,2,2,1,1,...,1,1,1,...,1,1,1,...,1),$$

$$\zeta((\alpha-1,2)|\Psi) = (1,2,1,1,1,...,2,1,1,...,1,1,1,...2),...,$$

$$\zeta((\alpha-1,\alpha-2)|\Psi) = (1,2,1,1,1,...,1,2,1,...,1,1,1,...1).$$

By the definition, we have attained unique metric representations for the elements of $G(\mathbb{Z}_{\alpha} \times \mathbb{Z}_{\alpha})$ Therefore, Ψ is a resolving set and $|\Psi| \le 3 + \alpha - 3 + \frac{\alpha - 3}{2} = \frac{3}{2}(\alpha - 1)$. **Theorem 3.6.** Let $\mathcal{R} \cong Z_{\alpha} \times Z_{\beta}$ where α and β are prime with $\alpha < \beta$, $\alpha = 3$, then $\beta(G(\mathcal{R})) \le \beta - 1$. *Proof.* Let $\Psi = \{(1,1), (2,2), (1,3), (2,4), (1,5), (2,6), ..., (1,\beta - 2), (2,\beta - 1)\}$.

Proof. Let
$$\Psi = \{(1,1), (2,2), (1,3), (2,4), (1,5), (2,6), \dots, (1,\beta-2), (2,\beta-1)\}.$$

In Fig.2, for any
$$(k,l)$$
, $(m,n) \in \mathcal{R}$, $d((k,l),(m,n)) = \begin{cases} 2, & \text{if } k+m \in U(\mathcal{R}) \text{ or } l+n \in U(\mathcal{R}) \\ 1, & \text{otherwise} \end{cases}$



The following metric representations are observed from Fig.2,



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\zeta((0,0)|\Psi) = (1,1,1,1,1,1,\dots,1,1),
                                                                                                    \zeta((0,1)|\Psi) = (1,1,1,1,1,1,\dots,1,2),
\zeta((0,2)|\Psi) = (1,1,1,1,1,1,\dots,2,1),
                                                                                                     \zeta((0,-1)|\Psi) = (2,1,1,1,1,1,\dots,1,1),
                                                                                                     \zeta((1,2)|\Psi) = (1,2,1,2,1,2,...,2,2),
\zeta((1,0)|\Psi) = (1,2,1,2,1,2,...,1,2),
\zeta((1,-1)|\Psi) = (2,2,1,2,1,2,...,1,2),
                                                                                                    \zeta((2,0)|\Psi) = (2,1,2,1,2,...,1,2,1)
                                                                                                    \zeta((2,-2)|\Psi) = (2,2,2,1,2,...,1,2,1)
\zeta((2,1)|\Psi) = (2,1,2,1,2,...,1,2,2),
By the definition, we have attained unique metric representations for the elements of (\mathbb{Z}_3 \times \mathbb{Z}). Therefore, \Psi is a
resolving set and |\Psi| \leq -1.
Theorem 3.7.Let \mathcal{R} \cong \mathbb{Z} \times \mathbb{Z} where and are prime with <, > 3, \ge 2-3, then (()) \le.
Proof. Let us denote \mathbb{Z}' = \mathbb{Z} \setminus \{0,1,-1\} = \{0,1,...,-4\}, \mathbb{Z}' = \mathbb{Z} \setminus \{0,1,-1\} = \{0,1,...,-4\}. Now define \Psi_1 = \{(0,1),(1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0),(-1,0
[1,-1], \Psi_2 = \{0,1,\ldots,-4\} where [1,-1], 0 \le \le -4. From the Construction of the above sets, \Psi_1 and \Psi_2 are
mutually disjoint to each other. Let \Psi = \Psi_1 \cup \Psi_2. The elements of \Re \Psi = UB can be written as follows.Let
 = \{(0,0), (1,1), (0,-1), (-1,0), (1,-1), (-1,1), (0), (0,), (1), (1,), (-1), (-1), (-1)\} where 0 \le \le -4, 0 \le \le -4
and \mathcal{B} = \{(a_{-3}), b\} where 0 \le a, t \le \beta - 4, s \ne t. We now prove that \Psi is a resolving set for G(\mathcal{R}).
d((0,0),(\alpha-1,\beta-1)) = 1, d((1,1),(\alpha-1,\beta-1)) = 2 \Longrightarrow \zeta((0,0)|\Psi) \neq \zeta((1,1)|\Psi).
d((0,\beta-1),(0,1)) = 2, d((\alpha-1,0),(0,1)) = 1 \Longrightarrow \zeta((0,\beta-1)|\Psi) \neq \zeta((\alpha-1,0)|\Psi).
d((1,\beta-1),(0,1)) = 2, d((\alpha-1,1),(0,1)) = 1 \Longrightarrow \zeta((1,\beta-1)|\Psi) \neq \zeta((\alpha-1,1)|\Psi).
d((0,\beta-1),(\alpha-1,\beta-1)) = 1, d((1,\beta-1),(\alpha-1,\beta-1)) = 2 \Longrightarrow \zeta((0,\beta-1)|\Psi) \neq \zeta((1,\beta-1)|\Psi).
Similarly, \zeta((0,\beta-1)|\Psi) \neq \zeta((\alpha-1,1)|\Psi), \zeta((\alpha-1,0)|\Psi) \neq \zeta((1,\beta-1)|\Psi),
\zeta((\alpha-1,0)|\Psi) \neq \zeta((\alpha-1,1)|\Psi).
d((0,0),(1,0)) = 2, d((0,\beta-1),(1,0)) = 1 \Longrightarrow \zeta((0,0)|\Psi) \neq \zeta((0,\beta-1)|\Psi).
Similarly, \zeta((0,0)|\Psi) \neq \zeta((\alpha-1,0)|\Psi).
d((0,0),(\alpha-1,\beta-1)) = 1, d((1,\beta-1),(\alpha-1,\beta-1)) = 2 \Longrightarrow \zeta((0,0)|\Psi) \neq \zeta((1,\beta-1)|\Psi).
Similarly, \zeta((0,0)|\Psi) \neq \zeta((\alpha-1,1)|\Psi).
d\big((1,1),(\alpha-1,\beta-1)\big)=2,d\big((0,\beta-1),(\alpha-1,\beta-1)\big)=1 \Longrightarrow \zeta\big((1,1)\big|\Psi\big)\neq \zeta\big((0,\beta-1)\big|\Psi\big).
Similarly, \zeta((1,1)|\Psi) \neq \zeta((\alpha-1,0)|\Psi).
d((1,1),(0,1)) = 1, d((1,\beta-1),(0,1)) = 2 \Rightarrow \zeta((1,1)|\Psi) \neq \zeta((1,\beta-1)|\Psi).
Similarly, \zeta((1,1)|\Psi) \neq \zeta((\alpha-1,1)|\Psi).
\forall (a_i, 0) \in \mathcal{R} \setminus \Psi and for any b_j \in \mathbb{Z}_\beta there exists (a_{\alpha - (i + \alpha - 3)}, b_j) \in \Psi such that d(a_i, 0), (a_{\alpha - (i + \alpha - 3)}, b_j) = 2 \Rightarrow 0
\zeta\big((a_i,0)\big|\Psi\big)\neq \zeta\big((a_l,0|\Psi)\big), 0\leq l\leq \alpha-4.
\forall (0, b_i) \in \mathcal{R} \setminus \Psi and for any a_i \in \mathbb{Z}_\alpha there exists (a_i, b_{\beta-(i+\alpha-3)}) \in \Psi such that
d\left((0,b_j),(a_i,b_{\beta-(j+\alpha-3)})\right)=2 \Longrightarrow \zeta\left((0,b_j)\big|\Psi\right) \neq \zeta\left((0,b_m)\big|\Psi\right), 0 \leq m \leq \beta-4,
\zeta((0,b_i)|\Psi) \neq \zeta((a_i,0)|\Psi),\zeta((1,b_i)|\Psi) \neq \zeta((1,a_m)|\Psi),0 \leq m \leq \beta - 4,
\zeta((a_i,1)|\Psi) \neq \zeta((a_i,1)|\Psi), 0 \leq l \leq \alpha - 4, \zeta((1,b_i)|\Psi) \neq \zeta((a_i,1)|\Psi).
d\big((a_i,0),(\alpha-1,\beta-1)\big)=1,d\big((a_i,1),(\alpha-1,\beta-1)\big)=2 \Longrightarrow \zeta\big((a_i,0)\big|\Psi\big)\neq \zeta\big((a_i,1)\big|\Psi\big).
Similarly, \zeta((0,b_i)|\Psi) \neq \zeta((1,b_i)|\Psi), \zeta((a_i,0)|\Psi) \neq \zeta((1,b_i)|\Psi), \zeta((0,b_i)|\Psi) \neq \zeta((a_i,1)|\Psi).
d\left((a_i,0),\left(a_{\alpha-(i+\alpha-3)},b_j\right)\right)=2,d\left((0,0),\left(a_{\alpha-(i+\alpha-3)},b_j\right)\right)=1 \Longrightarrow \zeta\left((0,0)\big|\Psi\right)\neq \zeta\left((a_i,0)\Psi\right).
Similarly, \zeta((0,0)|\Psi) \neq \zeta((0,b_i)|\Psi),\zeta((0,0)|\Psi) \neq \zeta((a_i,1)|\Psi),\zeta((0,0)|\Psi) \neq \zeta((1,b_i)|\Psi).
d\left((a_i,0),\left(a_{\alpha-(i+\alpha-3)},b_j\right)\right)=2,d\left((1,1),\left(a_{\alpha-(i+\alpha-3)},b_j\right)\right)=1 \Longrightarrow \zeta\big((a_i,0)\big|\Psi\big)\neq \zeta\big((1,1)\big|\Psi\big).
Similarly, \zeta\left(\left(0,b_{j}\right)\middle|\Psi\right)\neq\zeta\left(\left(1,1\right)\middle|\Psi\right),\zeta\left(\left(1,b_{j}\right)\middle|\Psi\right)\neq\zeta\left(\left(1,1\right)\middle|\Psi\right),\zeta\left(\left(a_{i},1\right)\middle|\Psi\right)\neq\zeta\left(\left(1,1\right)\middle|\Psi\right).
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d\left((a_i,0),(a_{\alpha-(i+\alpha-3)},b_i)\right)=2,d\left((\alpha-1,0),(a_{\alpha-(i+\alpha-3)},b_i)\right)=1 \Rightarrow \zeta((a_i,0)|\Psi)\neq \zeta((\alpha-1,0)|\Psi). Similarly,
\zeta((a_i,0)|\Psi) \neq \zeta((\alpha-1,1)|\Psi).
d((a_i, 0), (0, 1)) = 1, d((0, \beta - 1), (0, 1)) = 2 \Longrightarrow \zeta((a_i, 0)|\Psi) \neq \zeta((0, \beta - 1)|\Psi).
Similarly, \zeta((a_i, 0)|\Psi) \neq \zeta((1, \beta - 1)|\Psi), \zeta((0, b_i)|\Psi) \neq \zeta((\alpha - 1, 0)|\Psi),
\zeta\left(\left(0,b_{i}\right)\middle|\Psi\right)\neq\zeta\left(\left(\alpha-1,1\right)\middle|\Psi\right),\zeta\left(\left(0,b_{i}\right)\middle|\Psi\right)\neq\zeta\left(\left(0,\beta-1\right)\middle|\Psi\right),\zeta\left(\left(0,b_{i}\right)\middle|\Psi\right)\neq\zeta\left(\left(1,\beta-1\right)\middle|\Psi\right).
d\left((a_i,1),\left(a_{\alpha-(i+\alpha-3)},b_i\right)\right)=2,d\left((\alpha-1,0),\left(a_{\alpha-(i+\alpha-3)},b_i\right)\right)=1 \Rightarrow \zeta((a_i,1)|\Psi)\neq \zeta((\alpha-1,0|\Psi)).
Similarly, \zeta((a_i, 1)|\Psi) \neq \zeta((\alpha - 1, 1)|\Psi), \zeta((a_i, 1)|\Psi) \neq \zeta((0, \beta - 1)|\Psi),
\zeta((\alpha_i,1)|\Psi) \neq \zeta((1,\beta-1)|\Psi),\zeta((1,b_i)|\Psi) \neq \zeta((\alpha-1,0)|\Psi),\zeta((1,b_i)|\Psi) \neq \zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,0)|
\zeta((1,b_i)|\Psi) \neq \zeta((0,\beta-1)|\Psi),\zeta((1,b_i)|\Psi) \neq \zeta((1,\beta-1)|\Psi).
  \forall (a_i,\beta-1) \in \mathcal{R} \setminus \Psi \text{ and for any } b_i \in \mathbb{Z}_\beta \text{ there exists } \left(a_{\alpha-(i+\alpha-3)},b_i\right) \in \Psi \text{ such that } d\left((a_i,\beta-1),\left(a_{\alpha-(i+\alpha-3)},b_i\right)\right) = 0
2 \Longrightarrow \zeta \big( (a_i,\beta-1) \big| \Psi \big) \neq \zeta \big( (a_l,\beta-1) \big| \Psi \big), 0 \leq l \leq \alpha-4.
Similarly, \zeta((\alpha-1,b_j)|\Psi) \neq \zeta((\alpha-1,b_m)|\Psi), 0 \leq m \leq \beta-4.
d\left((a_i,\beta-1),\left(a_{\alpha-(i+\alpha-3)},b_j\right)\right)=2,d\left((\alpha-1,a_i),\left(a_{\alpha-(i+\alpha-3)},b_j\right)\right)=1 \Longrightarrow \zeta\left((a_i,\beta-1)\big|\Psi\right)\neq \zeta\left(\left(\alpha-1,b_i\right)\big|\Psi\right).
Similarly, \zeta((a_i, \beta - 1)|\Psi) \neq \zeta((0,0)|\Psi), \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((0,0)|\Psi), \zeta((a_i, \beta - 1)|\Psi) \neq \zeta((1,1)|\Psi), \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((0,0)|\Psi), \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1, b_i)|\Psi) \neq \zeta((\alpha - 1,
\zeta((1,1)|\Psi),\zeta((a_i,\beta-1)|\Psi) \neq \zeta((0,\beta-1)|\Psi),\zeta((\alpha-1,b_i)|\Psi) \neq \zeta((0,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi) \neq \zeta((0,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi) \neq \zeta((0,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi) \neq \zeta((0,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi) \neq \zeta((0,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi) \neq \zeta((0,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi) \neq \zeta((0,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi)) \neq \zeta((0,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi),\zeta((a_i,
\zeta((\alpha-1,0)|\Psi),\zeta((\alpha-1,b_i)|\Psi) \neq \zeta((\alpha-1,0)|\Psi),\zeta((a_i,\beta-1)|\Psi) \neq \zeta((1,\beta-1)|\Psi),\zeta((\alpha-1,b_i)|\Psi) \neq \zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi) \neq \zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi) \neq \zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),\zeta((\alpha-1,b_i)|\Psi),
\zeta((1,\beta-1)|\Psi),\zeta((a_i,\beta-1)|\Psi) \neq \zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,b_i)|\Psi) \neq \zeta((\alpha-1,1)|\Psi),\zeta((a_i,\alpha-1)|\Psi) \neq \zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha-1,1)|\Psi),\zeta((\alpha
\zeta((a_i,0)|\Psi),\zeta((\alpha-1,a_i)|\Psi) \neq \zeta((a_i,0)|\Psi),\zeta((a_i,\alpha-1)|\Psi) \neq \zeta((a_i,1)|\Psi),\zeta((\alpha-1,a_i)|\Psi) \neq \zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((a_i,1)|\Psi),\zeta((
\zeta((a_i,\alpha-1)|\Psi) \neq \zeta((0,a_i)|\Psi),\zeta((\alpha-1,a_i)|\Psi) \neq \zeta((0,a_i)|\Psi),\zeta((a_i,\alpha-1)|\Psi) \neq \zeta((1,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0,a_i)|\Psi),\zeta((0
\zeta((\alpha-1,a_i)|\Psi) \neq \zeta((1,a_i)|\Psi).
For any a \in \mathbb{Z}_a, d\left((a_i,b_j),(a,b_{\beta-(j+\alpha-3)})\right) = 2 \Longrightarrow \zeta\left((a_i,b_j)|\Psi\right) \neq \zeta((a_l,b_m)|\Psi) where
0 \le l \le \alpha - 4, 0 \le m \le \beta - 4.
Similarly, \zeta((a_i, b_j)|\Psi) \neq \zeta((0,0)|\Psi), \zeta((a_i, b_j)|\Psi) \neq \zeta((1,1)|\Psi), \zeta((a_i, b_j)|\Psi) \neq \zeta((0,\beta-1)|\Psi),
\zeta\left(\left(a_{i},b_{j}\right)\middle|\Psi\right)\neq\zeta\left(\left(\alpha-1,0\right)\middle|\Psi\right),\zeta\left(\left(a_{i},b_{j}\right)\middle|\Psi\right)\neq\zeta\left(\left(1,\beta-1\right)\middle|\Psi\right),\zeta\left(\left(a_{i},b_{j}\right)\middle|\Psi\right)\neq\zeta\left(\left(\alpha-1,1\right)\middle|\Psi\right).
d((a_i, b_i), (1,0)) = 1, d((a_i, 0), (1,0)) = 2 \Longrightarrow \zeta((a_i, b_i)|\Psi) \neq \zeta((a_i, 0)|\Psi).
  d((a_i, b_i), (0,1)) = 1, d((0, b_i), (0,1)) = 2 \Longrightarrow \zeta((a_i, b_i)|\Psi) \neq \zeta((0, b_i)|\Psi).
d\left((a_i,b_i),(\alpha-1,\beta-1)\right)=1,d\left((a_i,1),(\alpha-1,\beta-1)\right)=2 \Longrightarrow \zeta\left((a_i,b_i)|\Psi\right)\neq \zeta((a_i,1)|\Psi).
d\left((a_i,b_i),(\alpha-1,\beta-1)\right)=1,d\left((1,b_i),(\alpha-1,\beta-1)\right)=2 \Longrightarrow \zeta\left((a_i,b_i)|\Psi\right)\neq \zeta\left((1,b_i)|\Psi\right).
d((a_i,b_j),(0,1)) = 1, d((a_i,q-1),(0,1)) = 2 \Longrightarrow \zeta((a_i,b_j)|\Psi) \neq \zeta((a_i,q-1)|\Psi).
d((a_i,b_i),(1,0)) = 1, d((\alpha-1,b_i),(1,0)) = 2 \Longrightarrow \zeta((a_i,b_i)|\Psi) \neq \zeta((\alpha-1,b_i)|\Psi).
Hence \Psi is a resolving set for G(\mathcal{R}).
```

We now prove that removal of any element from Ψ does not result in resolving set.If $\Psi' = \Psi \setminus (0,1)$, then $\zeta((0,0)|\Psi) = \zeta((\alpha-1,0)|\Psi)$.Similarly, if $\Psi' = \Psi \setminus (1,0)$, then $\zeta((0,0)|\Psi) = \zeta((0,\beta-1)|\Psi)$.If $\Psi' = \Psi \setminus (\alpha-1,\beta-1)$, then $\zeta((0,\beta-1)|\Psi) = \zeta((1,\beta-1)|\Psi)$. If $\Psi' = \Psi \setminus (\alpha-1,\beta-1)$, then $\zeta((0,\beta-1)|\Psi) = \zeta((0,\beta-1)|\Psi)$ where $0 \le k \le \beta-4$. Therefore, Ψ is a resolving set and $|\Psi| \le \beta$.

CONCLUSION

For over a decade, there has been considerable research on the metric dimension of graphs. Metric dimension of zero divisor graph, total graph and other such graphs have been computed already. In this work, the metric dimension of unit graph is computed while the metric dimension of other algebraic graphs are up for research.

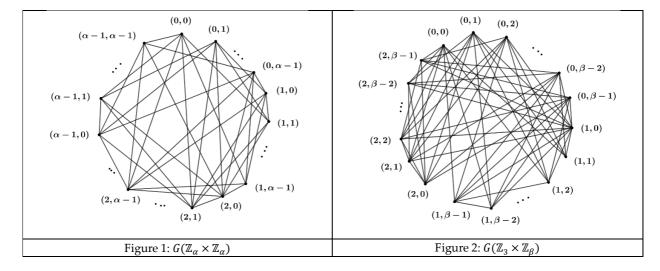




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RESEARCH ARTICLE

Predictive Modeling for Breast Cancer Detection: A Logistic Regression **Approach**

Imran Qureshi

College of Computing and Information Sciences, UTAS, Al Musanna Sultanate of Oman

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*Address for Correspondence Imran Qureshi

College of Computing and Information Sciences, UTAS, Al Musanna Sultanate of Oman Email: imranqureshi1210@gmail.com



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ABSTRACT

A pestilence disease that kills people in large numbers annually is breast cancer. On a global scale, it ranks as the leading cancer killer of females. When it comes to health-related information classification, ML approaches are the way to go. Classification and analysis make heavy use of it to help in the decisionmaking process. As in this study, analysis of logistical regression is one of the many ML classification models used to analyses the WBCD... The study aims to determine how good the classifiers performed in classification of the datasets. To achieve this, we used all the features in our dataset through the RFE approach. To evaluate the replica's efficiency, k-fold cross validation metric was used. All experiments were conducted using Anaconda Jupiter notebooks in Python language. Empirical analysis shows that the logistic regression model has a very high prediction accuracy of 96.7 percent for tumor benign or malignant in breast cancer Wisconsin.

Keywords: Breast Cancer; WBCD; classifiers, Logistic regression

INTRODUCTION

Cancer is a spectrum of illnesses signified by clonal expansion of rogue cells that can arise from any organ or tissue in the body [1]. In 2018, the disease was responsible for about 9.6 million deaths. This disease is characterized by its heterogeneity and can be classified into various unique forms. Breast cancer, a form of cancer, is the second most mutual malignancy in females, following carcinoma of the lung [2]. Finding out how well and efficiently the classifiers classified the dataset is the main goal of this investigation. We used the Recursive Feature Elimination method to select all of the dataset's characteristics for achieving this goal. We measured the model's efficacy using the K-fold cross-validation metric. We executed every single investigation in the Anaconda environment using Jupiter Notebooks and the Python computer programming language. In 2018, Nigeria recorded a total of 115,000 ongoing cancer cases, which led to over 70,000 fatalities attributed to the illness. In addition, breast cancer accounted





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for 22.7% of all cases, with a mortality rate of 16.4%. The breast cancer trend chart projected a significant rise in reported cases, estimating that by the year 2040, the number of active cases will nearly double from 26,310 in 2018 to around 50,921 [1]. Figure 1 illustrates a visual representation of breast cancer tumor sizes, with T1 representing tumors that are 2cm or smaller, and T4 representing tumors that have grown into the chest wall regardless of their size. Early identification of cancer increases the likelihood of effective therapy by 30%, whereas late discovery complicates the remedy process [4,5]. Medical practitioners apply many ways to detect preliminary carcinoma of the breast, involving surgical biopsy, which is approximately 100% accurate. [4], The Fine Needle Aspiration procedure involves visually interpreting the results, which have an accuracy rate ranging from 65% to 98% [6], and mammography with 63% to 97% accuracy [7]. Although it's expensive, the first method is the most dependable. The WBCD dataset will be evaluated in this study using ensemble ML classifiers based on FNA.

A form of AI called machine learning trains computers to acquire new skills by feeding them examples of information from actual real-world problems [8]. In recent decades, ML has been widely adopted to develop prediction models that allow better decision-making across different areas. This approach may be advantageous for the realm of cancer research due to which it can detect deviating patterns in datasets and utilize such unusual patterns in forecasts. To date, several papers have focused on enhancing computational analysis of FNA specimens over the last two decades [9]. To assess the WBCD dataset, this study employs a pair of ensemble machine learning classifiers: XG Boost and RF. In section 2, there is a summary of relevant studies. Section 3 provides an in-depth analysis of the cancer dataset utilized for the try out. The experimental procedure and the ideas of two ensemble machine learning classifiers that were studied are described in detail: section 4. In Section 5, the results and discussion of the experiment are given. This work contains sections 6 and, both of which offer advice, as well as a conclusion.

Related Work

Perhaps the most widely known application of machine learning is classification task. For example, many studies apply machine learning classifiers on various medical data sets such as WBCD dataset. Numerous studies have shown that the classifiers employed in those studies gave high classification accuracies. As one of the studies by Salama et al., their work was done under "Breast Cancer Diagnosis on Three Different Datasets Using Multi-Classifiers" using WEKA data mining system [10]. This study employed the WBCD dataset, one of the breast cancer data sets. Five types of AI classifications were used to do a binary classification job on the datasets: These are NB, MLP, J48 SMO and IBK. According to the results of these WBCD experiments, SMO classifier showed better performances with a mean classification accuracy of 94.5637 % and max value: 96.9957% as compared to IBK classification scheme Therefore, it was not difficult to yield an accuracy rate of 97.568 using J48 and M [11] have analyzed feature selection for breast cancer data from three separate datasets. In the study, binary classification was used when the CART classifier was adopted. That is, experiments showed nearly 94.84% classification rate for this classifier when it was used on WBCD without feature reduction. Lastly, after trying a variety of feature selection methods on the dataset, CART with exhaustive elimination produced a high accuracy. But the combination of these six descriptors results in 95.13% precision level. A set of machine learning classifiers such s Simple Logistic, RepTree and RBF Network were tested in [12]. The sample for this study was drawn from the Institute of Oncology, University Medical Center Ljubljana in Yugoslavia. The dataset has 286 rows and ten features. From the results of an experimental study, it is observable that Simple Logistic classifier yields better results in a binomial classification test with an accuracy percentage of 74.5%. Asri et al. [13] applied varied machine-learning algorithms in detecting and diagnosis of cancer risk for breast cancer. Performance evaluation was performed by comparing the performance of SVM, C4.5, NB and kNN algorithms with WBCD dataset for classification of cancer data correctly. The findings of the experimental results indicated highest level accuracy for SVMs had a nine percent error score rating as lowest. The research used the WEKA data mining tool. Therefore, I performed a review to assess and compare the power of machine learning algorithms used in breast cancer diagnosis. Three machine learning approaches utilised the analytical component of the study included support vector machines , random forests and Bayesian networks . For this experiment, the algorithms' performance was evaluated using WBCD dataset. The study used the WEKA data mining tool for simulation which indicated an accuracy rate of 97%. Second, the selected method may have a





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substantial effect on the outcome of categorization. Support Vector Machine (SVM) can be compared to the sensitivity, detail and accuracy. Although, random forest had the highest likelihood of classification on breast cancer data. Ivančáková et al. [15] conducted a comparison of diverse machine learning methods using the Wisconsin Dataset. The dataset used had 569 unique records with 32 attributes. Six machine learning classifiers were used to simulate the study experiment. The dataset was partitioned into three different ratios (60:For the purposes of assessing, 40%, 70:30 and 80:20 were selected. I did some research to assess and compare the competencies of machine learning algorithms in detecting and identifying breast cancer. The study presented three machine learning methods: SVM, RF, and BN. The experiment confirmed the efficiency of the algorithm applied by using WBCD database. As for the WEKA data mining tool that was applied by the study, the accuracy of 97% as a result in testing phase. Moreover, the chosen method can have significant influence on the classification result. As for reliability, sensitivity and precision, SVM has performed better. On the other hand, random forests had the biggest likelihood of correctly classifying the breast cancer dataset. As the utilization of data mining methods, a model was constructed from breast cancer data by Shajahaan et al. [16]. The WBCD data set was also used to try various machine learning classifiers such as RF, CART, C4.5, ID3 and NB. In addition, the processes of simulated data were carried out with TANGARA and WEKA programs associated with data mining. The highest accuracy was demonstrated by the NB classifier with 97.42% obtained from experiment results, which is relatively perfect in this study. The work Amrane et al. [17] runs kNN and NB as two ML classifiers aimed at the classification of breast cancer. WBCD data were used to assess the accuracy of classifiers. In simulation results, the KNN classifier obviously outperforms all others with an accuracy of 97.51% as compared to 96.19%. In terms of the total running time, the NB classifier was quicker in comparison with kNN model. In their paper, Bayrak et al. [18] examined the performance of two machine learning algorithms - Artificial Neural Network and Support Vector Machine (SVM), used in breast cancer diagnosis. Based on the WBCD dataset, strategy effectiveness was evaluated using WEKA data mining tool in the experiment. In case the sequential minimal optimization algorithm is implemented for the dataset to be diagnosed and predicted, its parameters are optimal with 96.9957% accuracy. Ahmad et al. [19] use three ML algorithms (C4.5, ANN, and SVM) to model breast cancer onset predictors. An Iranian ICBC dataset was used in this analysis. The dataset has 1,189 cases, with 22 predictors and one dependent variable. When the records went missing, 547 were left in the list after preprocessing. On the accuracy point of view, SVM classifier performed better (95.7%) compare to C4.5's 93.6% and ANN had 94.7%. We performed all experiments using the only WEKA data mining tools. Lastly, with a view to discriminate and diagnose the levels of risk for breast cancer through machine learning algorithms, Showrov et al. [20 took several approaches such as C4.5, ANN and SVM that were used in order to predict if a patient will develop or not breast cancer. This study used a data set from the ICBC in Iran. Overall, the dataset had 1,189 total observations; of these there were 22 predictors and one was an outcome variable. At the conclusion of the pre-treatment stage, only 547 records were left due to missing data regarding the issues that disrupted a total number of 642 entries. In contrast to the C4.5 and ANN algorithms, our SVM algorithm yielded a better performance score of 95.7 compared to 93.6% while using binary tasks with two classifiers namely Naive Bayes and k-Nearest Neighbours. This performance from the experiment showed that kNN classifier accuracy was on 97.51% and NB got a high rating of 96.19%. On the contrary, scientists also revealed that while kNN was more accurate, its compute time would rise if there were a larger set of data in the experiment. The accuracy of C4.5 classifier in classification was measured by Sarkar and Nag [22] using the WBCD dataset. The classifier was used to solve a binary classification problem. For the classifier, experimental results were 96.71% accuracy. Java 7 programming language was used for the studies. Using C4.5 decision tree-based categorization systems, the study's authors found that their performance improved early stage detection in individuals with risk factors for breast cancer. This might also possibly save thousands of lives annually. Houfani and his colleagues did a study titled "Breast cancer classification using machine learning techniques: a comparative work [23], based on the WBCD database. The datasets were binary-classified using seven different artificial intelligence classifiers: SVM, RF, DT, MLP, LR, NB and kNN. According to the results of the experiment, classification accuracy varied from 95% for NB to 97.9% for MLP and LR classifiers. Among the classifiers tested, MLP and LR performed well. Two ensemble classifiers that are used in this study are RF and EGB. For the classifier's performance evaluation, we adopted RFE technique to reduce features in the dataset. There are two key areas where machine learning algorithms learn: supervised learning and unsupervised learning. In this case,





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the issue is supervised learning that requires a dataset of labeled data examples in order to train the system so as it produces correct outcomes (classification). In this study we selected supervised learning classifiers.

MATERIALS AND METHODS

There are 699 occurrences in the breast cancer Wisconsin dataset, as stated in the publication. Ten characteristics make up the breast cancer dataset in Wisconsin. Among the characteristics included in the Wisconsin breast cancer dataset are the following: sample code number, lump thickness, cell size uniformity, cell shape consistency, and marginal adhesion. Dimensions of a single epithelial cell, Standard nucleoli, nuclei without chromatin, Mitoses Table 1: Summary of the Dataset

METHODOLOGY

The suggested research approach consists of the following procedures: The breast cancer dataset from Wisconsin was obtained by gathering information using the UCI machine learning repository. [24]. Information preprocessing is used to handle missing values and prepare the dataset for logistical regression model training. The modeling process uses Python for logistic regression. Separating the dataset into a training set and a test set allows one to train a logistic regression model on the former and then assess its efficacy on the latter. K-fold cross-validation provides a more robust metric for evaluating the model's correctness and effectiveness on fresh data. Logistic regression, a statistical tool, aids in binary classification by estimating the likelihood of a binary result. Logistic regression allows one to make certain conclusions about the type of tumor, benign or malignant, from the Wisconsin data set on breast cancer provided some conditions are met. The logistic regression model is a derivation of the logistic function or the sigmoid function that indicates how input features correspond to the probability of binary outcome. The estimated probability must lie within the bounds of 0 and 1 for the logistic function to be valid. The model generates the log odds of the binary result by using coefficients to represent input variables. The model estimates the propensity towards binary result with logistic function based on log-odds. Logistic regression is often employed for training a model on labeled data, assessing the quality of the model and then using it to make forecasts on new instances. It is possible to assess the performance of a model using different measures such as accuracy, precision, recall and area under the ROC curve. But logistic regression is a great option when you need to figure out the relationship between input data and binary output, for it produces clear results that are popular in binary classification projects. As its mathematical representation, logistic regression makes use of the logistic function, which is

 $P(Y=1 \mid X) = \frac{1}{1 + e^{-(\frac{beta_0 + beta_1X_1 + beta_2X_2 + ... + beta_pX_p)}}$ (1)

P(Y=1|X)) is the probability of the binary outcome being 1 given the input features X. e is the base of the natural logarithm.

\beta_0, \beta_1, \beta_2, ..., \beta_p are the coefficients estimated by the model.

 $X_1, X_2, ..., X_p$ are the input features.

A logistic function can be used for binary classification because it guarantees that the predicted probabilities range from 0 to 1. Given that logistic regression uses coefficients (\beta_0, \beta_1, \beta_2,..., \beta_p) to depict the influence of input features on the estimated binary outcome rate.

Data Preprocessing

We used Jupyter Notebook, an integrated development environment (IDE) for Python, to generate the classifiers. We chose this option because Python is a robust programming platform that has been regularly used in previous studies of this kind.. Python offers numerous built-in libraries that are commonly utilised during the dataset pre-processing stage. The initial step undertaken was the importation of the dataset into the graphical user interface (GUI) in order to comprehend and visually represent the dataset. The results showed that 16 cancer reports had missing or partial information; thus, they were removed. The experiment utilized the remaining 683 data points. More so, the sample code number element was eliminated as it contains the patient's identifying number, which is superfluous for





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classification. In addition, there are two parts to the experiment; the first part makes use of every characteristic in the dataset. Step two included applying Recursive Feature Elimination (RFE) to the dataset in order to reduce the number of features. The feature selection approach determined that five features were the best fit for the categorization problem. Among these traits are mitoses, normal nucleoli, uniform cell size and shape, and bland chromatin. In addition, we used the min-max normalization approach to rescale the features to values between 0 and 1, which helped reduce the dataset's intrinsic skewness. The following formula describes it [24].

 $Y=x-\min(x)/\max(x)-\min(x)$

Classifier Evaluation

For the evaluation of this logistic regression model, we used a technique termed K-fold cross validation. To assess the logistic regression model, we used K-fold cross-validation. Likewise, a confusion matrix was adopted to assess the precision of correct and wrong guesses on test set. With these assessment frameworks, we evaluated the model generalizability outside of training information and its performance on novel observations. It has good accuracy to the extent that a logistic regression model can predict benign or malignant tumors in breast cancer Wisconsin dataset approximately up to 96.7%. The k-fold cross validation metric used to evaluate the proposed model was deemed to be a more reliable measure of accuracy, and the results therefore indicate that its variance in performance was moderate across different test folds. Physicians and organizations could evaluate whether the logistic regression model is of practical utility in predicting outcomes for women with breast cancer. These conclusions suggest that the logistic regression model has implementation capacity into forecasting tumor characteristics on which health care and medical decision-making can be built. Our model on the logistic regression model demonstrates a high level of accuracy, approximately 96.7%, in predicting benign or malignant tumors in the breast cancer Wisconsin data set.

Future Scope & Recommendations

But its efficacy or not could have been found out from other medical data sets and hence, its wide applicability to medicine also may be evaluated. Illuminating more features or creating new ones in the dataset could improve the prediction power of models and provide a better insight into the tumor properties. This model performance assessment used a method known as K-fold cross-validation. To assess the performance of the logistic regression model, we used K-fold cross-validation. We also investigated the accuracy and misjudgment of predictions on test set with confusion matrix. We evaluated the extrapolation ability of the model with these testing protocols, also enabling us to estimate its performance on new data.

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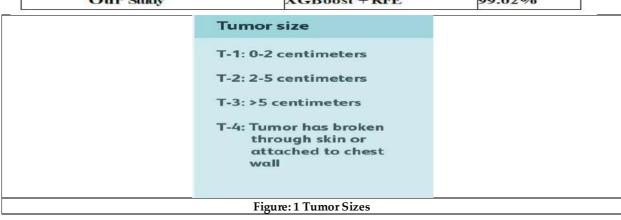
Imran Qureshi

Table 1: Summary of the Dataset

S/N	Attribute	Domain	
1	Sample code number	id number	
2	Clump Thickness	1-10	
3	Uniformity of Cell Size	1-10	
4	Uniformity of Cell Shape	1-10	
5	Marginal Adhesion	1-10	
6	Single Epithelial Cell Size	1-10	
7	Bare Nuclei	1-10	
8	Bland Chromatin	1-10	
9	Normal Nucleoli	1-10	
10	Mitoses	1–10	
11	Class	2 for Benign & 4 for Malignant	
Class Distribution		Benign: 458 (65.5%); Malignant: 241 (34.5%)	
Total N	umber of All Instances	699	
Numb	er of Missing Values	16	

Table 2: Papers using WBCD.

Ref	Classifiers	Accuracy
Lavanya and Rani [11]	CART + Exhaustive (FST)	95.13%
Chaurasia and Pal [12]	Simple Logistic	74.5%
Asri and his colleagues [13]	SVM	97.13
Bazazeh and Shubair [14]	SVM	97%
Shajahaan and his colleagues [16]	NB	97.42
Ahmad and his colleagues [19]	SVM	95.7%
Showrov and his colleagues [20]	SVM	96.72%
Amrane and his colleagues [21]	kNN	97.51%
Sarkar and Nag [22]	C4.5	96.71%
Houfani and his colleagues [23]	MLP & LR	97.9%
Sumbaly and his colleagues [30]	J48	94.36%
Bennett and Blue [31]	SVM	97.20%
Chaurasia and his colleagues [32]	NB	97.36%
Islam and his colleagues [33]	ANN	98.57%
Our Study	XGBoost + RFE	99.02%







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RESEARCH ARTICLE

Modelling Threshold Logic Neuron as a Linear Filter for Pattern Classification

S. Arulraj¹ and M. Subbulakshmi. N. Murugan²

¹Department of Mathematics, St. Xavier's College (Autonomous), Palayamkottai 627 002, Tamil Nadu, India.

²Research Scholar (Reg.No:20111282092006), PG and Research Department of Mathematics, St. Xavier's College (Autonomous), Palayamkottai 627 002, Tamil Nadu, India.

^{1,2}Affiliated to Manonmaniam Sundaranar University, Abishekapatti, Tirunelveli 627 012, Tamil Nadu, India.

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*Address for Correspondence

S. Arulraj

Department of Mathematics, St. Xavier's College (Autonomous), Palayamkottai 627 002, Tamil Nadu, India. Email: arulrajsj@gmail.com



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ABSTRACT

For the past few years, the thirst to explore the human brain and to design a machine to mimic human intelligence has paved the way for neural network models to spread their application in various domains of engineering. In this artificial intelligence-driven culture, neural networks play a significant role in solving the pattern classification problem. A literature review revealed that binary neurons are pattern dichotomizers in two-dimensional Boolean space. In this paper, we have extended the two-dimensional Boolean space to n-dimensional linearly separable Boolean space and implemented Threshold Logic Neuron as a pattern classifier. In the present context, the Threshold parameter and the weight vector have been fixed, and the binary threshold signal function is used as the neuronal activation function. Threshold Logic Neuron is employed to approximate a hyperplane, which acts as a tool to classify the given linear separable pattern set into two distinct classes.

Keywords: Threshold Logic Neuron, Pattern set, Threshold parameter, neuronal activation function, hyperplane.





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INTRODUCTION

The father of the modern brain [7] referred to neurons as the mysterious butterflies of the soul, by beating its wings could provide insights into the secret of mental life. The organization of the human brain is far too complex to analyse and quantify in numeric terms. The history of neuroscience dates to the period of Greek civilization which sown the seed about the structure and function of the human brain [7]. The thirst to design an engineering system that can endure the Turing test [7] and model the complex structure of the human brain is ever ending process. The field of learning has expanded its horizon from humans to machines. In this 21st century machines do learn, make errors, and make corrections thanks to the advancement in the field of artificial intelligence. In [1] McCulloch and Pitts perceived that the brain consists of entities with logic-gating potential. The origin of the neural network is motivated by the information-processing entity of the brain called the neuron. In 1943 McCulloch and Pitts gave birth to the concept of the "Boolean brain" [7] and introduced the first neural network to form a mathematical model. The artificial neural network model [9] is represented as a mathematical model of a natural neural network. Artificial neural networks can be applied as a problem-solving tool in various fields like classification, pattern recognition, and optimization. Pattern recognition [2] involves the study of the process by which machines can learn from the environment and differentiate patterns of concern to predict appropriate and make rational decisions about the pattern classes. According to [6]CNNs extract features better than traditional algorithms in image classification, object detection, and speech recognition. [5] proposed a new CNN architecture for classification.A CNN was designed for pneumonia detection, using transfer learning and pattern recognition. [8] ANN can predict supercritical fluid heat transfer accurately in thermodynamics. [4] B. C. Geigerhas thrown light on the issue of estimating mutual information in deterministic NNs and addressed its consequences. In [7] Mc Culloch and Pitts proposed a highly simplified computational model of the brain, which is referred to as MP neuron and is used for pattern classification in 2D linearly separable Boolean space. In the present context, MP neuron is referred as Threshold Logic Neuron (TLN) with binary threshold signal function. Motivated by [7] we extend the two-dimensional linearly Boolean space to n-dimensional linearly separable Boolean space and employ TLN as a two-class pattern classifier.TLN is also used to approximate the hyperplane which is the decision surface to classify the linearly separable pattern set into two classes.

Preliminaries

The following notation discussed below are employed in this paper. $X = (x_1, x_2, ..., x_n)^T$ here X denotes the vector and $x_1, x_2, ..., x_n$ denotes its component and T refers to the transpose of the vectors. The χ is used to denote the pattern set taken into consideration.

Definition[7]

A Boolean $f(x_1, x_2, ..., x_n)$: $B^n \to \{0, 1\}$ is linearly separable, if there exist a hyperplane π in \mathbb{R}^n that strictly separates χ_0 from χ_1 and $\pi \cap \{0, 1\}^n = \emptyset$.

Architecture of TLN

The TLN consist of two parts. The first part receives the input signal x_i from various sources. These input signal travels through a weighted path w_i and neuronal activation a_l is generated, where l is the index of the TLN taken into consideration. The internal activation a_l is linear weighted summation of the input vectors, modified by the internal threshold parameter θ_l . That is,

$$a_l = \sum_{i=1}^n w_{i_l} x_i \tag{1}$$

The activation a_l represent the inner product of the input vector $X = (x_1, x_2, ..., x_n)^T$ and weight vector $W_l = (w_1, w_2, ..., w_n)$ $a_l = X^T W_l = \sum_{i=0}^n w_{i_l} x_i$





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(2)

Where $x_0 = +1$ and $w_0 = -\theta_l$.

In the second part, the neuronal activation a_l represented in (2) is now transferred to a signal function $\delta(.)$, which is

$$\delta(a_l) = \begin{cases} 0 & \text{if } a_l < 0 \\ 1 & \text{if } a_l > 0 \end{cases}$$
 (3)

defined as follows:

In this study signal function $\delta(a_l)$, is Binary threshold signal function, which yields the output $y_l = 1$ for positive neuronal activity and yields output $y_l = 0$ for negative neuronal activity.

Threshold Parameter (θ)

Consider the neuronal activation of TLN represented in the equation (2)

$$a_{l} = \sum_{i=0}^{n} w_{i_{l}} x_{i}$$

$$= \sum_{i=1}^{n} w_{i_{l}} x_{i} + w_{0} x_{0}$$

$$= m_{l} + \theta_{l}$$

Where θ_l is the threshold parameter of TLN modelled by $w_0 = -\theta_l$ and $x_0 = 1$. Here, m_l refers to the neuronal activation from external sources of input. And θ_l is threshold internal to the neuron. For the θ_l consider the following cases.

Case 1: $\theta_1 = 0$

In this case the net activation of the neuron is from external input sources. That is a_l refers to neuronal activation, when threshold is set to zero. That is the signal function $\delta(.)$ is always centered at 0. For neuronal activational value in the range $(-\infty, 0)$, the signal function yield output $y_l = 0$ and for the range $(0, \infty)$ it yields output $y_l = 1$.

Case 2: $\theta_1 = k \neq 0$

In this case, the threshold parameter θ_{l} , biases the neuron to yield output $y_{l} = 0$ for the net external inputs in range $(-\infty, +k)$, and yield output $y_{l} = 1$ for the net external inputs in range $(+k, \infty)$.

Measures of Activation

The activation a_l represented in equation (2) is simply the inner product of input vector $X = (x_0, x_1, ..., x_n)^T$ with the weight vector $W = (w_0, w_1, ..., w_n)^T$. That is, $a_l = X^T W_l$

$$= \sum_{i=0}^{n} w_{i} x_{i} \text{ where } w_{0} = -\theta_{l} \& x_{0} = 1$$
 (4)

This inner product depicts the similarities or dissimilarities measure between the input vector and the weight vector. For larger neuronal activation, the similarity between the input vector and weight vector is large. Otherwise, it yields a relatively smaller neuronal activation which implies the input vector is dissimilar to weight vector. We can interpret the TLN as Linear filter, as neuronal activation measures help to filter the input vector that are similar or dissimilar to weight vector respectively.

Problem Formulation

Space of Input pattern setAn n-dimensional Boolean function is a map that assign each n-dimensional vector in domain space B^n into unique element in the set $\{0,1\}$.

$$f \colon B^n \to \{0,1\} \tag{5}$$

The function given in (5) is a Boolean function.





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For the present study, we restrict linearly separable Boolean function and domain B_n to its finite subset $\chi = \{X_1, X_2, ..., X_p\}$, where χ is the input pattern set under consideration.

For the input pattern $\chi \subseteq B^n$, the corresponding Boolean function

$$f_{\chi}(x_1, x_2, ..., x_n) : B^n \to \{0, 1\}$$
 (6)

is defined such that.

$$\chi_0 = f_{\chi}^{-1}(0) = \{(x_1, x_2, \dots, x_n) \in \chi | \forall x_i \neq 1; 1 \le i \le n\}$$

$$\chi_1 = f_{\chi}^{-1}(1) = \{(x_1, x_2, \dots, x_n) \in \chi | \forall x_i = 1; 1 \le i \le n\}$$

This paper aims to classify the pattern set χ into two class χ_0 and χ_1 using TLN.

TLN: As Two Class Pattern Classifier

Now let us proceed further to approximate a hyperplane, π in \mathbb{R}^n that separates χ_0 and χ_1 and $\pi \cap \chi = \emptyset$ using TLN which is illustrated below.

Let $\theta = \frac{2n-1}{2}$ be the threshold parameters for the TLN. Let $W = (w_1, w_2, ..., w_n)$ and $w_i = 1 \forall i = 1, 2, ..., n$. The pattern set $\chi = \chi_0 \cup \chi_1$ is the input pattern set of vectors to the threshold logic neuron (TLN) respectively. Now, let $\chi_0 = \chi_0' \cup \chi_0'$ where.

$$\chi_0 = \{(x_1, x_2, \dots, x_n) \in \chi_0 | x_i = 0; \ \forall 1 \le i \le n\}$$

$$\chi_0 = \{(x_1, x_2, \dots, x_n) \in \chi_0 | \ \forall x_i \ne 0; \ \forall 1 \le i \le n\}$$

Let $X_j \in \chi_0$ which implies $X_j \in \chi_0$ or $X_j \in \chi_0$. The suffix j is the index of the input vector taken into consideration.

Case 1: Let $X_i \in \chi_0$ be the input vector to the TLN. Let

$$\sum_{i=1}^{n} w_i x_{ji} = a_j \tag{7}$$

Where a_i is the inner product of the j^{th} input vector X_i^T with the weight vector W.

Therefore $a_i = 0$. Let

$$b_j = a_j - \theta \tag{8}$$

$$b_j = 0 - \frac{2n-1}{2} < 0$$

Now, the activation b_j in equation (7) for the input vector X_j received from the aggregation part of the neuron is send to the Binary threshold signal function δ where

$$\delta(b_j) = \begin{cases} 1 & \text{if } b_j > 0 \\ 0 & \text{if } b_j < 0 \end{cases} \tag{9}$$

$$\delta(b_i) = y_i = 0 \tag{10}$$

Where y_i is the output of input vector X_i .

Case 2: Let $X_j \in \chi_0^{''}$ be the input vector to the TLN. That is,

$$\sum_{i=1}^{n} w_i x_{ji} = a_j \tag{11}$$

Therefore $a_i \le n-1$ since $X_i \in \chi_0^n$. Consider, For,

$$b_i = a_i - \theta$$

$$b_j = n - 1 - \frac{2n - 1}{2} \le 0$$

Now, the activation b_j for the j^{th} input X_j received from aggregation part of the TLN sent to the Binary threshold signal function δ , which yields final output y_j .

$$\delta(b_j) = 0 \qquad (\because b_j < 0)$$

i.e., $y_j = 0$



(12)



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Thus equation (10) and (12) suffices that for $X_j \in \chi_0$. The TLN yield output $y_j = 0$. Let $X_j \in \chi_1$ be the input vector to the TLN and

$$a_j = \sum_{i=1}^n w_i x_{ij}$$

Hence, we get that $a_i = n$ since all the component of X_i input vector is 1. Now for,

$$b_j = n - \theta$$

$$= n - \frac{(2n-1)}{2}$$

$$= 0.5 > 0$$

The activation b_j for the j^{th} input vector X_j is send to the Binary threshold signal function δ which yields the final output.

$$\ddot{\delta}(b_j) = 1 \ (\because b_j > 0)$$

$$i.e. \delta(b_j) = y_j = 1$$

$$(14)$$

From (10), (12) and (14) we observe that the weighted Σ of inputs yields a neuronal activation. That is, $y = X^T W = w_1 x_1 + w_2 x_2 + \dots + w_0 x_0$

Where $w_0 = -\theta$, that is the threshold parameter θ is modelled by the weight w_0 . The function y(X) = 0 is called the discriminant function of the neuron that is it is the equation of the required hyperplane π defined in equation,

$$x_1 + x_2 + \dots - \frac{2n-1}{2} = 0 \tag{15}$$

Thus, TLN approximated the hyperplane π in \mathbb{R}^n given in the equation (15) classifies the pattern set, χ into χ_0 and χ_1 that is, $\forall X \in \chi_0$ on one side of the plane and $\forall X \in \chi_1$ on the other side respectively with $\pi \cap \chi = \emptyset$. Thus, we have illustrated that TLN is a two-class pattern classifier in n-dimensional linearly separable Boolean space.

CONCLUSION

In the present paper, we have implemented Threshold Logic Neuron as a linear filter and classified the n-dimensional linearly separable Boolean pattern space into two class pattern spaces. This study can be extended to the linear classification of multi-class patterns. Also, multilayer network of neurons could be employed to solve pattern classification problems.

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RESEARCH ARTICLE

Edge Sum Divisor Cordial Labeling of Crown & Comb Graphs

A. Anto Cathrin Aanisha^{1,2} and R. Manoharan³

¹Sathyabama Institute of Science and Technology, Chennai, India

²Department of Education, DMI-St. John The Baptist University, Mangochi, Malawi

³Department of Mathematics, Sathyabama Institute of Science and Technology, Chennai, India

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*Address for Correspondence

A. Anto Cathrin Aanisha

Sathyabama Institute of Science and Technology, Chennai, India Department of Education, DMI-St. John The Baptist University,

Mangochi, Malawi

Email: antocathrinaanisha@gmail.com



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ABSTRACT

Let $\Omega = (W(\Omega), E(\Omega))$ be a graph which is having neither loop nor multiple edges, where $W(\Omega)$ represent node set and $E(\Omega)$ represent line set and let h: $E(\Omega) \to \{1, 2, \dots \mid E(\Omega) \mid \}$ be a bijection. For each node u, give it a label of 1 if $2 \mid h(b_1) + h(b_2) + ... + h(b_s)$ and 0 if it doesn't where $b_1, b_2, ... b_s$ are edges that are incident with the node u. If the difference between nodes categorized 0 and 1 is less than or equal to 1, the function h is called ESDC labelling. A ESDC graph is one that has the ESDC labeling. In this paper, we prove that the crown graph Ct O K1 and the comb graph Pt O K1 ESDC graph are edge sum divisor cordial graphs when t is even.

Keywords: SDC graph, ESDC graph, Crown graph, Comb graph

INTRODUCTION

This paper only considers nontrivial and undirected graph. Graph labeling is one of the part of graph theory Graph labeling has direct application in a number of fields such as coding theory, the research of X-ray crystallography, communication networks and many more (Parthiban & Sharma, 2020). For the past 60 years, graph labeling has found popularity in the field of graph theory (Daisy et al., 2022). The effective connection between number theory and the structure of graphs is facilitated by graph labelling. Graph labeling is a mathematical concept that is concerned with assigning positive integers to nodes or lines or both. Vertex labelling is when the subject of labelling is a set of vertices. Edge labelling is the process of putting labels on a set of edges. And labelling is called "total labelling" if the





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domain has both vertices and edges (Raheem et al., 2022). The cordial labelling is one of the graph's labelling options. As an extension of cordial labeling, prime cordial labelling, total cordial labelling, Fibonacci cordial labelling, etc have come into being (Raheem et al., 2022). DC labeling was initiated by (Varatharajan et al., 2011). Many graphs are divisor cordial graphs. Especially (Barasara & Thakkar, 2022) proved some ladder related graphs are divisor cordial graph. The practice of DC labelling has been expanded to SDC labelling, DDC labelling, PD - divisor cordial labelling, and other similar practices. The ESDC labeling is the topic of discussion in our paper. This discussion continues from where the Sum divisor cordial labelling left off. SDC labeling was initiated by (Lourdusamy & Patrick, 2016). After that he proved some graph which are satisfied the axioms of sum divisor cordial graph. Like some star and bistar related graphs are SDC graphs (Lourdusamy & Patrick, 2019). It has been demonstrated by a great number of authors that many graphs are SDC graph. And (Raheem et al., 2022) examined disconnected graphs, including disjoint union of paths and subdivided star SDC graphs. Recently (Adalja & Ghodasara, 2018) proved splitting graph, shadow graph, sunlet graph, shell graph are SDC graphs. Not all graphs are pleasant SDC graphs. Investigating sum divisor cordial labelling for the graphs or families of graphs that permit sum divisor cordial labelling is both very intriguing and challenging(Lourdusamy & Patrick, 2016).SDC graph is extended to ESDC graph. ESDC labeling was introduced by (Vijayalakshmi et al., n.d.). Already they proved that path, cycle, fan, wheel, closed helm, flower graph and friendship graphs are ESDC graph. In this paper, we prove that the crown graph C_t O K₁ and the comb graph P_t O K₁ ESDC graph are edge sum divisor cordial graphs when t is even.

Definition: 1.1

Consider $\Omega = (W(\Omega), E(\Omega))$ be a graph which is having neither loop nor multiple lines, where $W(\Omega)$ represent node set and $E(\Omega)$ represent line set and let $h: W(\Omega) \to \{1, 2, \dots, |W(\Omega)|\}$ be a bijection. For each line x, give it a label of 1 if h(x) divides h(y) or h(y) divides h(x) and 0 if not. If the difference between lines categorized 0 and 1 is less than or equal to 1, the function h is called DC labelling. A DC graph is one that has the DC labeling (Varatharajan *et al.*, 2011). **Definition of SDC graph: 1.2**

Let $\Omega = (W(\Omega), E(\Omega))$ be a graph which is having neither loop nor multiple lines, where $W(\Omega)$ represent node set and $E(\Omega)$ represent line set and let h: $W(\Omega) \to \{1, 2, \dots, |W(\Omega)|\}$ be a bijection. For each line x, give it a label of 1 if 2/(h(x)+h(y)) and 0 if not. If the difference between lines categorized 0 and 1 is less than or equal to 1, the function h is called SDC labelling. A SDC graph is one that has the SDC labeling(Lourdusamy & Patrick, 2016).

Definition of EDSC graph: 1.3

Let $\Omega = (W(\Omega), E(\Omega))$ be a graph which is having neither loop nor multiple lines, where $W(\Omega)$ represent node set and $E(\Omega)$ represent line set and let $h: W(\Omega) \to \{1, 2, \dots |W(\Omega)|\}$ be a bijection. For each node u, give it a label of 1 if $2 \mid h(b_1) + h(b_2) + \dots + h(b_s)$ and 0 if it doesn't where $b_1, b_2, \dots b_s$ are edges that are incident with the vertex u. If the difference between nodes categorized 0 and 1 is less than or equal to 1, the function h is called ESDC labelling. A ESDC graph is one that has the ESDC labelling (Vijayalakshmi $et\ al.$, n.d.).

Definition of crown graph: 1.4

The crown Ct Θ Kt is the graph formed by linking each vertex of a cycle with a pendant edge (Lourdusamy & Patrick, 2016).

Definition of comb graph: 1.5

The comb $P_t \Theta$ K_1 is the graph formed by adding a pendant edge to each vertex of a path(Lourdusamy & Patrick, 2016).

MAIN RESULT

Theorem: 2.1

The crown graph $C_t \Theta K_1$ is an edge sum divisor cordial graph when t is even.

Proof:

Let Ω = $C_t \circ K_1$ where t is even.

Let $W(\Omega) = \{u_x; 1 \le x \le t \text{ and } v_x; 1 \le x \le t\}$ be a node set and $E(\Omega) = \{e_1, e_2, \dots e_{2t}\}$ be a line set, where $e_x = u_{x+1}, v_{x+1}; 1 \le x \le t-1$





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 $e_t = v_1 u_1$

 $e_{t+x} = v_x v_{x+1}$; $1 \le x \le t-1$

 $e_{2t} = v_t v_1$

Also $|W(\Omega)| = 2t$ and $|E(\Omega)| = 2t$

Define h: $E(\Omega) \rightarrow \{1, 2, \dots \mid E(\Omega) \mid \}$ as follows:

 $h(e_{t+2x-1})=4x-3, 1 \le x \le \frac{t}{2}$

 $h(e_{2x-1}) = 4x-2, 1 \le x \le \frac{t}{2}$

 $h(e_{t+2x}) = 4x, 1 \le x \le \frac{t}{2}$

 $h(e_{2x}) = 4x-1, 1 \le x \le \frac{t}{2}$

Then induced vertex labels are

 $h^*(v_{2x-1})=1, 1 \le x \le \frac{t}{2};$

 $h^*(v_{2x}) = 0, 1 \le x \le \frac{t}{2};$

 $h^*(u_{2x-1}) = 0, 1 \le x \le \frac{t}{2}$

 $h^*(u_{2x}) = 1, 1 \le x \le \frac{t}{2}$

Considering the previously described labelling pattern, we have

 $v_h(0)=v_h(1)=t$

Thus, $|v_h(0)-v_h(1)|=|t-t|=0\le 1$

Here $v_h(1)$ represents the number of nodes labeled with 1 and $v_h(0)$ represents the number of nodes labeled with 0 among all nodes in Ω .

Therefore, the crown graph $C_t O K_1$ is an edge sum divisor cordial graph when t is even.

Example: 2.2

The crown graph $C_t \Theta K_1$, in which t=6 is shown in figure (a).

From figure (a), $|v_h(0)-v_h(1)| = |6-6| = 0 \le 1$.

So, we conclude that the crown graph $C_t \circ K_1$, where t=6 is having ESDC labeling.

Hence the crown graph C_t ° K₁, in which t=6 is an ESDC graph.

Example: 2.3

The crown graph C_t O K₁, in which t=4 is shown n figure (b).

From figure (b), $|v_h(0)-v_h(1)| = |4-4| = 0 \le 1$.

So, we conclude that the crown graph $C_t \circ K_1$, where t=4 is having ESDC labeling.

Hence the crown graph C_t ^o K₁, in which t=4 is an ESDC graph.

Theorem: 2.4

The comb graph $P_t\Theta$ K_1 is an edge sum divisor cordial graph when t is even.

Proof:

Let Ω = $P_t \Theta$ K_1 be the comb graph when t is even.

Let $W(\Omega) = \{u_x; 1 \le x \le t \text{ and } v_x; 1 \le x \le t\}$ be node set and $E(\Omega) = \{e_1, e_2, \dots e_{2t-1}\}$ be a line set, where

Let $e_x = u_x v_x$; $1 \le x \le t$

and $e_{t+x} = v_x v_{x+1}$; $1 \le x \le t-1$

Also, $|V(\Omega)| = 2t$ and $|E(\Omega)| = 2t-1$

Define h: $E(\Omega) \rightarrow \{1, 2, \ldots \mid E(\Omega) \mid \}$

 $h(e_x) = t-1+x; 1 \le x \le t$

 $h(e_{t+x}) = x$; $1 \le x \le t-1$

Then induced vertex labels are

 $h^*(v_{2x-1}) = 0$; $1 \le x \le \frac{t}{2}$

 $h^*(v_{2x}) = 1$; $1 \le x \le \frac{t}{2}$

 $h^*(u_{2x-1}) = 1; 1 \le x \le \frac{t}{2}$

 $h^*(u_{2x}) = 0$; $1 \le x \le \frac{t}{2}$

Considering the previously described labelling pattern, we have





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 $v_h(0)=v_h(1)=t$

Here $v_h(0)$ represents the number of nodes labeled with 0 and $v_h(1)$ represents the number of nodes with 1 in Ω .

Thus, $|v_h(0)-v_h(1)|=|t-t|=0\le 1$

Hence comb graph $P_t O$ K_1 is edge sum divisor cordial labeling where t is even.

Example: 2.5

The comb graph $P_t O K_1$, in which t=4 is shown in figure (c).

From figure (c), $|v_h(0)-v_h(1)| = |4-4| = 0 \le 1$.

So, we conclude that the comb graph PtO K1, where t= 4 is having ESDC labeling.

Hence the comb graph $P_t \odot K_1$, in which t=4 is an ESDC graph.

Example: 2.6

The comb graph $P_t O K_1$, in which t=6 is shown in figure (d).

From figure (d), $|v_h(0)-v_h(1)| = |6-6| = 0 \le 1$.

So, we conclude that the comb graph $P_t O K_1$, where t=6 is having ESDC labeling.

Hence the comb graph Pt O K1, in which t=6 is an ESDC graph.

CONCLUSION

In this paper, we have shown that the crown graph C_t $^{\circ}$ K_1 and comb graph $P_t \Theta$ K_1 are an edge sum divisor cordial graph for t is even. Future research will also determine whether the crown graph C_t $^{\circ}$ K_1 and the comb graph $P_t \Theta$ K_1 are an edge sum divisor cordial graph or not when t is odd.

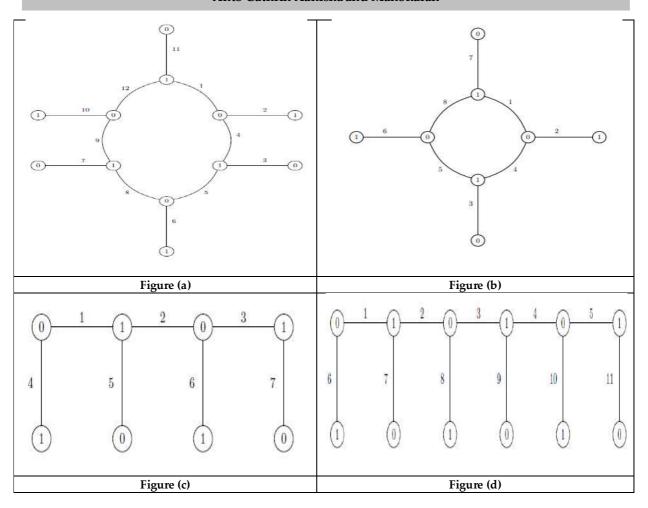
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RESEARCH ARTICLE

Enhancing Data Science Resources Within The Academic Library

P.J. Rosario Vasantha Kumar¹

Librarian, St. Xavier's College (Autonomous), Palayamkottai- 627 002

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*Address for Correspondence P.J. Rosario Vasantha Kumar

Librarian,

St. Xavier's College (Autonomous),

Palayamkottai- 627 002

Email: rosario.kumar@gmail.com



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ABSTRACT

The rise of data science as a pivotal interdisciplinary field has necessitated significant transformations within academic institutions, particularly in the resources and support provided by academic libraries. This research article explores the multifaceted role of academic libraries in enhancing data science resources and support within the university setting. Through an in-depth analysis of current practices, challenges, and opportunities, this article provides insights into the evolving landscape of data science within academic libraries. Drawing on literature review, case studies, and expert interviews, this article presents a comprehensive framework for optimizing data science resources and services within academic libraries. Key areas of focus include access to datasets, training and workshops, support for data management, access to software and tools, collaboration and partnerships, and research support services. By adopting a holistic approach to data science support, academic libraries can effectively meet the diverse needs of students, faculty, and researchers, and contribute to the advancement of data-driven research and innovation.

Keywords: Data Science, Academic Library, Resources, Support, Datasets, Training, Data Management, Software, Collaboration, Research Support

INTRODUCTION

In recent years, the exponential growth of data and the emergence of sophisticated analytical techniques have revolutionized various industries, research domains, and decision-making processes. The interdisciplinary field of





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data science, which integrates statistics, computer science, domain expertise, and communication skills, has become increasingly central to addressing complex societal challenges and driving innovation across diverse sectors. As the demand for data-driven insights continues to soar, academic institutions are recognizing the importance of fostering a robust data science ecosystem to equip students, faculty, and researchers with the requisite skills and resources. Academic libraries, as vital repositories of knowledge and information within universities, play a pivotal role in supporting data science education, research, and practice. Traditionally, libraries have served as custodians of scholarly resources, providing access to books, journals, and databases. However, with the advent of data science, libraries are adapting their services and collections to meet the evolving needs of the academic community. This research article seeks to explore the multifaceted role of academic libraries in enhancing data science resources and support within the university setting.

LITERATURE REVIEW

The interdisciplinary realm of data science is swiftly emerging as one of the most dynamic areas of exploration both within academia and beyond its borders. Integrating principles from statistics and computer science, data science is propelling groundbreaking discoveries across various fields. Examples abound, including automated linguistic analysis facilitating timely interventions in online support forums (Kornfield *et al.*, 2018), neuroimage processing predicting health outcomes (Lancaster, Lorenz, Leech, & Cole, 2018), and the study of environmental factors' impact on childhood cognitive development (Stingone, Pandey, Claudio, & Pandey, 2017). However, delving into such applications necessitates specialized skills and resources, encompassing computer programming and data visualization, which are indispensable for today's workforce (National Academies of Sciences, Engineering, and Medicine, 2018; Ridsdale *et al.*, 2015).

Regrettably, many academic institutions lack comprehensive data science support, resulting in a growing roster of unmet needs (Barone, Williams, & Micklos, 2017; Galanek & Brooks, 2018; Garcia-Milian, Hersey, Vukmirovic, &Duprilot, 2018; Oliver, 2017), thereby hindering researchers from fully capitalizing on the big data revolution. Several factors contribute to this gap in skills and resources vital for harnessing data science applications. Firstly, while data science as a field has existed for decades (Donoho, 2017), its widespread recognition across academic domains is a recent phenomenon. Consequently, many researchers are just now realizing the imperative of skill development (Barone *et al.*, 2017), often while holding positions—such as post-doctoral or faculty roles—where formal training proves especially challenging.

Engaging academic libraries in supporting data science endeavors presents a promising avenue for enhancing campus-wide data literacy (Martin, 2016; Maxwell, Norton, & Wu, 2018; Wang, 2013). Given their role as central hubs on college and university campuses, libraries naturally embody interdisciplinary collaboration, mirroring the ethos of data science. Moreover, the audience for academic libraries arguably surpasses that of any other campus unit, serving not only students but also staff and faculty. While the idea of libraries supporting aspects of data science is not novel—libraries have long championed data collection, management, and sharing best practices (Antell, Foote, Turner, & Shults, 2014)—some have also provided researchers with training in computational and statistical skills (e.g., North Carolina State University: https://www.lib.ncsu.edu/services/data-visualization, New York University Libraries: https://library.nyu.edu/departments/data-services/, Arizona State University: https://lib.asu.edu/data). Thus, the current landscape presents a ripe opportunity for academic libraries to explicitly address the growing demand for data science training.





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Engaging academic libraries in supporting data science endeavors presents a promising avenue for enhancing campus-wide data literacy (Martin, 2016; Maxwell, Norton, & Wu, 2018; Wang, 2013). Given their role as central hubs on college and university campuses, libraries naturally embody interdisciplinary collaboration, mirroring the ethos of data science. Moreover, the audience for academic libraries arguably surpasses that of any other campus unit, serving not only students but also staff and faculty. While the idea of libraries supporting aspects of data science is not novel—libraries have long championed data collection, management, and sharing best practices (Antell, Foote, Turner, & Shults, 2014)—some have also provided researchers with training in computational and statistical skills (e.g., North Carolina State University: https://www.lib.ncsu.edu/services/data-visualization, New York University Libraries: https://library.nyu.edu/departments/data-services/, Arizona State University: https://lib.asu.edu/data). Thus, the current landscape presents a ripe opportunity for academic libraries to explicitly address the growing demand for data science training.

Access to Datasets

One of the primary roles of academic libraries in supporting data science is providing access to a diverse array of datasets for research purposes. Libraries acquire and curate datasets from various sources, including government agencies, research institutions, and commercial providers. These datasets cover a wide range of domains, including social sciences, natural sciences, healthcare, economics, and more. By offering access to datasets, libraries facilitate interdisciplinary research and foster innovation across diverse fields.

Training and Workshops

Many academic libraries offer training sessions, workshops, and tutorials on data science tools, techniques, and methodologies. These sessions are designed to help students, faculty, and researchers develop essential data literacy skills, such as data manipulation, visualization, statistical analysis, and machine learning. Moreover, libraries often collaborate with campus units, such as computer science departments, statistics departments, and interdisciplinary research centers, to deliver comprehensive training programs tailored to the needs of the academic community.

Data Management Support

Effective data management is essential for conducting robust and reproducible research. Academic libraries provide guidance and support to researchers in managing, organizing, and sharing their data in compliance with best practices and institutional policies. This includes assistance with data storage, metadata creation, data documentation, version control, and data preservation strategies. Moreover, libraries play a key role in advocating for open data initiatives and promoting data sharing and transparency within the research community.

Access to Software and Tools

Libraries often license and provide access to a variety of data science software and tools, including statistical analysis software, programming languages, data visualization platforms, and machine learning libraries. By offering these resources, libraries ensure that students and researchers have the necessary tools to analyze and interpret data effectively. Moreover, libraries may provide training and support services for these tools, helping users navigate complex software environments and leverage advanced functionalities.

Collaboration and Partnerships

Academic libraries collaborate with other campus units, such as research centers, computer science departments, and interdisciplinary programs, to strengthen data science initiatives. By forging partnerships and leveraging expertise





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from diverse areas, libraries enhance their ability to support cutting-edge research and innovation in data science. Moreover, libraries may collaborate with external stakeholders, such as industry partners, government agencies, and non-profit organizations, to access additional resources, expertise, and funding opportunities for data science initiatives.

Research Support Services

Libraries offer a wide range of research support services tailored to the needs of data scientists and researchers. This includes assistance with literature searches, citation management, research impact analysis, data citation, and research data management planning. Librarians with expertise in data science and information retrieval play a pivotal role in guiding researchers through the complex landscape of scholarly communication and data discovery. Moreover, libraries may provide consultation services, workshops, and training sessions on research methodologies, data analysis techniques, and scholarly publishing practices. The impact of data science on academic libraries is profound and multifaceted, influencing various aspects of research, education, and scholarly communication. By embracing data science initiatives, academic libraries can effectively address the evolving needs of researchers, students, and faculty, ultimately enhancing the quality and impact of scholarly endeavors. Here are some key ways in which data science initiatives impact academic libraries:

The Impact Of Data Science On Academic Libraries

Facilitating Research Innovation: Data science initiatives enable academic libraries to support innovative research projects by providing access to diverse datasets, advanced analytical tools, and expertise in data analysis methodologies. This facilitates interdisciplinary collaboration and empowers researchers to explore complex research questions, leading to novel discoveries and insights across disciplines.

Enhancing Teaching and Learning Academic libraries play a crucial role in promoting data literacy and computational skills among students through workshops, training programs, and access to educational resources. By incorporating data science concepts into curriculum development, libraries help students develop critical skills for navigating the data-rich landscape and conducting rigorous research.

Promoting Open Science and Data Sharing Academic libraries advocate for open science principles by promoting data sharing, transparency, and reproducibility in research. By providing platforms for hosting and disseminating research data, libraries contribute to the advancement of open access initiatives and foster collaboration among researchers worldwide.

Supporting Scholarly Communication Data science initiatives in academic libraries enhance scholarly communication by enabling researchers to leverage data visualization techniques, interactive dashboards, and text mining tools to communicate their findings more effectively. This facilitates knowledge dissemination and encourages interdisciplinary collaboration within the academic community.

Bridging the Digital Divide Academic libraries play a crucial role in bridging the digital divide by providing equitable access to data science resources and support for students and researchers from diverse backgrounds. By promoting inclusivity and accessibility, libraries ensure that all members of the academic community have the opportunity to engage in data-driven research and education. Overall, the impact of data science initiatives in academic libraries is transformative, empowering researchers, students, and faculty to harness the power of data for discovery, innovation, and scholarly advancement. By embracing data science initiatives, academic libraries can fulfill their mission of supporting research, teaching, and learning in the digital age, ultimately contributing to the advancement of knowledge and society as a whole.

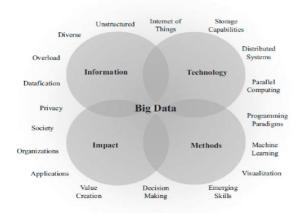




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Challenges in Strengthening Data Science Resources within Academic Libraries

Enhancing data science resources within the academic library presents various challenges that must be addressed to effectively meet the evolving needs of researchers, students, and faculty.



These Challenges Include

Lack of Specialized Expertise Academic libraries may face challenges in recruiting and retaining staff with specialized expertise in data science methodologies, tools, and technologies. Building a team of skilled professionals capable of providing comprehensive support for data science initiatives can be difficult, especially in competitive job markets.

Limited Funding and Resources Acquiring and maintaining data science resources, such as access to datasets, software licenses, and computational infrastructure, requires significant financial investment. Academic libraries may struggle to secure adequate funding to procure these resources, especially in the face of competing priorities and budget constraints.

Technological Complexity Data science tools and technologies are constantly evolving, requiring library staff to stay abreast of new developments and trends. Keeping up with the rapid pace of technological change can be challenging, particularly for libraries with limited technical expertise and resources. Data Management and Governance: Academic libraries must navigate complex data management and governance issues, including data privacy, security, and compliance with regulatory requirements. Ensuring the responsible and ethical use of data while also facilitating access and sharing presents a delicate balancing act for library staff.

Interdisciplinary Collaboration Data science is inherently interdisciplinary, requiring collaboration between researchers from diverse fields. Academic libraries must foster partnerships with other campus units, research centers, and external stakeholders to effectively support interdisciplinary data science initiatives. Overcoming institutional silos and promoting collaboration across departments can be challenging but is essential for success.

Promoting Data Literacy: Promoting data literacy among students, faculty, and researchers is essential for leveraging data science resources effectively. Academic libraries must develop and deliver data literacy programs, workshops, and educational resources to empower users with the skills and knowledge needed to navigate the data-rich landscape.

Infrastructure and Access Issues Ensuring equitable access to data science resources and infrastructure can be challenging, particularly for institutions with limited physical or technological resources. Addressing issues of accessibility, usability, and inclusivity is essential for ensuring that all members of the academic community can benefit from data science initiatives.





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Resistance to Change Embracing data science initiatives may require cultural and organizational changes within academic libraries, including shifts in priorities, workflows, and decision-making processes. Overcoming resistance to change and fostering a culture of innovation and experimentation is essential for driving progress in enhancing data science resources within the academic library. Addressing these challenges requires strategic planning, collaboration, and investment in staff training and development. By overcoming these obstacles, academic libraries can effectively enhance their data science resources and support, ultimately advancing research, teaching, and learning within the academic community.

CONCLUSION

The integration of data science resources within academic libraries represents a significant leap forward in meeting the evolving needs of researchers, students, and faculty within the university setting. The transformative impact of data science initiatives on academic libraries is evident across various fronts, including research innovation, teaching and learning enhancement, promotion of open science, scholarly communication support, and bridging the digital divide. By embracing data science, libraries are not only adapting to the changing landscape of scholarly communication but also actively contributing to the advancement of knowledge and society as a whole. However, this journey toward enhancing data science resources within academic libraries is not without its challenges. From the necessity for specialized expertise to limited funding and resources, technological complexity, data management concerns, interdisciplinary collaboration barriers, and resistance to change, libraries face multifaceted obstacles that require strategic planning and concerted efforts to overcome. Yet, by addressing these challenges through collaboration, investment in staff training, and fostering a culture of innovation, libraries can effectively navigate the complexities of data science and continue to serve as vital hubs for research, teaching, and learning in the digital age. Moving forward, it is imperative for academic libraries to remain agile and proactive in adapting to emerging trends and technologies in data science. By staying abreast of developments, forging partnerships, and advocating for the principles of open science and data literacy, libraries can play a pivotal role in empowering the academic community to leverage the power of data for discovery, innovation, and societal impact. Through these collective efforts, academic libraries will continue to fulfill their mission of supporting research, education, and scholarly communication in an increasingly data-driven world.

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RESEARCH ARTICLE

A Role of Artificial Intelligence In Smart Building

AR.A.Bhuvaneshwari^{1*} and A.Natarajan²

¹Professor, Department of Architecture, Sigma College of Architecture, Kuzhithurai-629168, Tamil Nadu. ² Assistant Professor, PG Department of Data Science, St. Xavier's College, Palayamkottai-627002, Tamil Nadu, India.

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*Address for Correspondence AR. A. Bhuvaneshwari

Prof/Dept Of Architecture, Sigma College Of Architecture, Kuzhithurai, Tamil Nadu 629168 Email: natrajanbhuvana@gmail.com



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ABSTRACT

Smart building uses artificial intelligence in various stages of construction of a building like designing, material selection, construction management, structures and maintenance. In this paper case study of five smart buildings of different countries using artificial intelligence has been analysed. The study reveals the usage of architectural intelligence to enhance design, project management, safety, and sustainability. With the evolution of artificial intelligence we can expect more innovations and efficiency in the building industry.

Keywords: Smart Buildings, Artificial Intelligence, Energy Efficiency.

INTRODUCTION

Artificial intelligence has made revolution in building industry in optimising designs, project management, predicting hazards, and optimising energy use. Artificial intelligence helps in making cost effective, safe and sustainable designs. It makes eco-friendly choices to attain sustainability.

Artificial Intelligence

Artificial intelligence is the intelligence of machines or software, as opposed to the intelligence of other living beings, primarily of humans. It is a field of study in computer science that develops and studies intelligent machines.

Smart Building

Smart buildings take advantage of many existing technologies and are designed or redesigned to allow for the integration of future technological developments. Internet of things (iot) sensors, building management, artificial intelligence (ai) and augmented reality are some of the techniques and technologies that can be used in these smart





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homes to manage and improve their performance. Current standards for smart homes cover areas such as health, energy and security. this model can operate continuously through relationships with users or self-management. Smart home solutions should not only be about the relationship between buildings and their users, but also about the outside and places like smart cities and smart communities. Smart buildings are important for the low-cost economy in the future, and the use of information technology and smart buildings will be efficient and integrated. In the long term, smart home implementation will require multiple layers of software/hardware adaptability to achieve high-quality services and reduce energy/cost major improvements require new hardware, understanding of all system issues, and integration.

Study Of Role Of Ai In Smart Buildings Through Case Studies

The Shanghai Tower, China The Edge, Amsterdam The One Central Park, Sydney The Biqbuilding, Hamburg. The Museum Of The Future, Dubai.

Case Study1

The Shanghai Tower, China (2015)

Architect: Gensler

Structural Engineer: Thornton Tomasetti

Mep: Cosentini

Fire Protection And Life Safety: Rja Fire Protection Technology Consulting (Rja) Building Façade: Aurecon Engineer Consulting (Shanghai) Co., Ltd. (Aurecon)

About Shangai Towers

Shangai world financial centre is the tallest building of china with a china green building three star rating. The shanghai tower is also a leed gold-rated building . It is the second tallest building of the world after burj khalifa in dubai.

AI Applications In Shangai Towers

The shanghai is the best example for usage of artificial intelligence in design and construction of sky scrapers. Few such applications are as follows.

Energy Efficiency

AI algorithm was used to study the building occupancy pattern and weather data to control the heat, cool and light of the building. It reduces the energy consumption of the building.

Structural Stability

AI was used to study the wind patterns as it was an important treat for structural stability. It created a unique, twisting form to reduce wind loads.

Selection Of Materials

AI was used to select material in the basis of structural stability and environmental impact. This helped with the sustainable principles of the tower.

Project Management

Artificial Intelligence was used to schedule construction activities, allocate resources efficiently, and monitor progress. It helped in achieving time and financial management.

Safety Measures

AI is used for sensing hazards and saving workers. They used sensors and cameras.





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Customization

Ai was used in designing the interior space according to the tenants need. Ai was used in complete customization.

Case Study2

The Edge, Amsterdam(2015)

Architect :Plp Architects Structural Engineer:Van Rossum Mep:Deerns Building Façade:Rollecate

About The Edge

The edge is an office building in amsterdam. It provides a better environment with sustainable properties. It is the world's highest breeam rating awarded to an office building.

AI Applications In Edge

The edge has been admired as the smartest building of the world. It has been rightly called as the computer with a roof. Its artificial intelligence application is as follows

Smart Office

Artificial intelligence suggests workers desk location based on temperature preference and meeting locations. On days when there are less employees expected some section might be shut down to reduce the cost of heating, cooling, lighting and cleaning. Artificial intelligence informs when to fill the coffee machine. It even informs when to clean up a busy bathroom, robots are used for office security.

Smart Parking

When we enter the garage a camera snap a photo of the license plate. It checks with the employees record and raised the gate. Sensor light brighten when one approaches and dims as we leave.

Case Study 3

The One Central Park, Sydney, (2014)

Architect: Ateliers Jean Nouvel

Structural Engineer: Robert Bird Group

Mep: Arup

Building Façade: Surface Design Pty Ltd

About The One Central Park

The one central park is a living architectural project in Australia. The building is visually appealing greenery in urban context.

AI Applications In One Central Park

The one central park has five green star rating. Artificial intelligence has been used for designing its greenery.

Green Facade

Artificial intelligence was used to face the challenges due to microclimate. Planting design and technology used were tested for long term planting success. For the façade wind and shade analysis were done to analyze the suitability of plant.in foyer and atrium lux level is recorded to determine low level light plant.

Case Study 4

The Biq Building, Hamburg, (2014):

Architect :splitterwerk





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Bio energy façade: arup

About The Biqbuilding, Hamburg

The biq building is a four floor residential building, it is the first bio energy façade building, the project won the zumtobel group award 2014.

AI Applications In Biq Building

The biq building has 129 glass panels to cultivate microalgae. Ai usage in this building is as follows.

Algae Growth

The residential building is made of algae bricks that produces oxygen and absorbs carbon dioxide. Ai used to control the building ventilation system and optimize the algae growth.

Case Study 5

The Museum Of Future, Dubai (2022)

Architectural Firm: Killa Design Structural Engineer: Burahappold

Exterior Arabic Calligraphy: Emirati Artist Mattar Bin Lahej

About Future Museum

The museum of the future's has a unique form that represents image of the future. It has received leed platinum certification.

AI Applications In Future Museum

The calligraphy and the building's unusual appearance make it one of the most complex architectural projects ever completed. Artificial intelligence was used in the construction of the building.

Design Of The Building

If you don't have the skills, parametric design and design knowledge this will be an impossible task. Parametric design is a technique based on algorithmic reasoning that allows certain variables or parameters to change the outcome of the equation. Bim is a 3d model-based technology for construction professionals to collaboratively design and document projects.

Comparison Of Case Studies

Show Table 1

CONCLUSION

Ai using different algorithm produces multiple design options based on material selection and construction cost. This helps architects to explore multiple design considering key features like strength, sustainability and aesthetics. It makes revolution in the field of architecture by streamlining decision making and designing more efficient and sustainable buildings.

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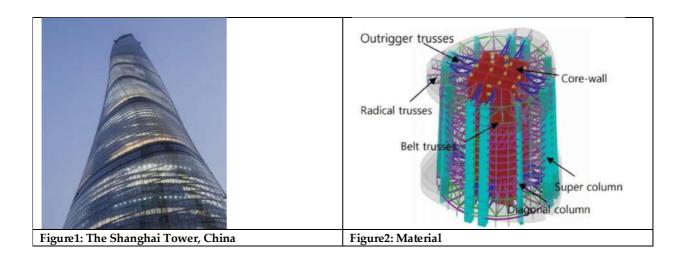


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Table1: Comparison Of Case Studies

Example	Location	Type Of	Ai Usage	
		Building		
Shangai Towers	China	Mixed Use	Energy Efficiency, Structural Stability,	
			Selection Of Materials, Project Management,	
			Safety Measures, Customization	
The Edge	Amsterdam	Office	Smart Office, Smart Parking	
The One Central Park	Sydney	Mixed Use	Green Facade	
The Biq Building	Hamburg	Residential	Algae Growth	
The Museum Of	Dubai	Public	Design Of The Building.	
Future				
Interference	Ai Is Used In Building Industry In Various Stages Right From Designing, Material			
Selection, Construction, Maintenance And Operation.				







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Figure 3: The Edge, Amsterdam







Figure 5: Biq Building





RESEARCH ARTICLE

LTE: Performance Investigation for The Mode of Interfacing in The **Integrated Hetnets**

¹Burhanuddin Mohammad and ²Imran Qureshi

^{1,2}Lecturer, College of Computing and Information Sciences-IT Department, University of Technology and Applied Science-AlMusanna, Sultanate of Oman

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*Address for Correspondence Burhanuddin Mohammad

Lecturer,

College of Computing and Information Sciences-IT Department, University of Technology and Applied Science-Al Musanna, Sultanate of Oman

Email: burhanuddin@act.edu.om

<u>@0</u>99

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ABSTRACT

Networking techniques are advancing quickly especially in case of pertaining to wireless communication systems. Researchers are looking forward to make the wireless technology as the future of networking. Since the evolution of wireless technologies rapid proliferation has been observed in wireless technologies. Simultaneously heterogeneity in technological development is observed. This heterogeneity comes up due to the application for the specific group of users. Now the demand with the development of the various wireless technologies is to bring all the heterogeneous technologies under one Umbrella i.e. Long-Term Evolution. LTE is a key development in wireless technologies. And LTE is considering the heterogeneous wireless technologies to integrate them. This research paper proposed the possible research challenges while integrating the different wireless technologies. And we consider the two scenarios for designing the framework tight and loose coupling. We simulated the both scenarios and observed how well the system works. Our research indicates that Tight coupling is better than the loose coupling.

Keywords: WiMax, LTE, HetNets, WLAN, Integration of wireless networks.

INTRODUCTION

Various wireless technologies are emerged and became obsolete before implementations few are very good case in point are Adhoc networks, and measuring device networks. Such innovations have valuable societal implications and are restricted to particular consumers. The evolution of LTE will have a significant impact on the incorporation of such innovations. For the simple reason that it is compatible with packet switched innovations and allows high-





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data-rate hybridized networking. In recent years WiMax (IEEE802.16e) gained great heed due to facilitate massive amounts of information assigns, high standards of services and support to movement and wider coverage area [6]. The 3rd Generation Partnership Project most presently specified the global LTE to meet the increasing working requirements of wireless internet access in cellphones . An attractive possibility for making the 4G paradigm a reality is the cooperation and amalgamation of multiple innovations [7]. Because the protocols and procedures for supporting quality of service in LTE connections are distinct. Integrating the innovations alongside the help of internetworking needs standard adaption. If we were to use layer two strategy, for instance, the The WiMax network BS and LTE eN would need to have their medium access control layers adapted. An LTE user will only ever interface with their respective LTE serving gateway (S-GW) under a layer 3 architecture, where adaption is executed at the IP layer. This WiMax/LTE integrated network is best suited for it 3 technique. No changes to the WiMax base station or LTE user devices are necessary because an LTE S-GW is in charge of protocol adaption up to the IP layer [5]. And at this level integration and internetwork of Adhoc and wireless sensor network has no obligation because WiMax, Adhoc and Sensor network are already supporting the same routing protocols like AODV, OLSR, TORA and GRP.

Advancements in such modular design enabling consumers to modify swiftly among the heterogeneous networks would provide numerous beneficial to both end users and operators. By providing comprehensive LTE/WiMax/Adhoc/WSN services, users would be benefitted from the increased performance and tremendous data rate of such integrated services and unlimited applications. For now, the WiMax subscriber owns the necessary LTE connectivity infrastructure but not restricted and can be owned by any other party. For this integration and ownership . For seamless cooperation, appropriate guidelines and terms of service must be established [5]. Now it is the perfect moment to consider how to incorporate diverse technological innovations into the LTE network. In addition, we need to zero in on the problems and obstacles associated with doing investigation into the complete portable network's cooperation, execution, and activation. Wireless connectivity provide a number of significant investigation issues. There will be a plethora of new problems to solve as we attempt to include differing wireless innovations, which will inevitably bring their own set of difficulties. The subsequent investigation obstacles require attention and conversation.

Research Challenges

Framework for Integration

A seamless switching across different types of communication networks is a crucial need for the next generation wireless system. Radio access networks have different features and technologies to provide the services to the user. UMTS and WiMax provide the support for a scarce data transport capacity that allows for greater accessibility. WLAN supports the less degree of mobility with high data transmission bandwidth. And for coverage a large number of access points needed. A temporary network allows for a great deal of getting around, lacks a centralised design, and has a geometry that is constantly changing to accommodate different levels of data transfer. Variegated degrees of accessibility are accommodated via Zigbee or smart sensor networks, which need minimal data transfer and provide concentrated assistance. The standards and limits that allow different types of organizations to deliver their capabilities to users are well specified. Therefore, linking all of the disparate WiFi networks is an enormous undertaking that calls for an extremely complex foundational architecture. Several ports and couplings are covered here to facilitate the creation of an LTE foundation and the integration of various wireless networks. When planning a melding arrangement, it is possible to think about

Open Interface

Independent connections and transportation routes were required by the open communication. Even though it's the most basic interface, it's rather complicated to come together. It incorporates a significant postponement over changeover between sites and requires an additional authentication method. If we stick to the accessible interface, we'll end up with inadequate connectivity and data delays during transfer, which goes against the whole point of LTE. This means it should be reserved for use as a last resort and main adoption [5].





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Tight/Close Interface

The principle of a network controller is derived from the notion of snug or intimate interconnection. During a transition, the link's access point that connects both systems acts as a radio network controller. Here, many radio technologies function as an integrated fundamental infrastructure. Substantial data transmission rates and smooth transitioning between connections are the foundation of LTE, and although the interface is a bit complicated, it provides these features [5].

Loose Interface

A disintegrating connection connects the WLAN network's AAA server to the WiMax framework using a shared authorization process. So, instead of modifying the WLAN and WiMax standards, users can have accessibility to capabilities through a loose integration.

Mobile IP based Interface

In order to facilitate the world of technology portability of points through the process of packet swapping idea, mobile IP is a top option for implementing the network connection. The care of address from the person's residence connection is used to identify cellphone users [8]. When moving clients on the move across different networks, mobile IP is the easiest and most effective way to do it missing affecting the functionality [9].

Clients can easily move across networks with the use of cell-phone IP [10]. However, information packet flipping across diverse networks is supported via packet wrapping in a standard arrangement [11].

Layered Protocols

Intensive and thorough investigation is required to develop a methodology as the foundation for Hspa-based networking in order to preserve the core of LTE and framework reliability. Due to the fact that various architectural styles employ distinct groups of methods. Investigators should prioritize the implementation of regulations for LTE-based combined networks.

Channel Allocation

Regulated broad range is being utilized by WiMax as well as other network operators. gratis levels are utilized by WLAN, Adhoc, WSN, and Zigbee connections. Channel allocation conflicts are a natural consequence of technology interconnection. Furthermore, it will generate a significant amount of distraction. A number of transmit placement strategies are at your disposal, including hybrid approaches, dynamic, and regular channels.

Handoff

Vertical handoff between between two different networks and horizontal handoff between the two different access points within the same network is available.

Security

Integrating diverse networks and LTE-based systems poses serious threats to security. Tradeoffs exist within safety and latency. We must find a way to solve this challenge better than everyone else. Completely coordinated safety features for networks are notoriously difficult to create all at once. However, we may go step-by-step. If you want your data exchange to be both quick and trustworthy, you may add layers of protection [36, 37, 38, 39, and 40].

Quality of Service

Quality of service concerns are less pressing when technologies are in their early stages of advancement. However, we must ensure that the system's fundamental performance remains intact. The standard of service may be categorized into two groups: those based on internal networks and those based on global networks [27], [28], and [29], respectively. The reason being that various networks offer varying levels of assistance, and the minimum requirement worldwide for excellence performance ought to match the highest level of the internally network's QoS.

Related Work

The goal of the future-oriented infrastructure is to standardize the building blocks and foundation for all portable and connected systems, regardless of their heterogeneity. Access to the network made available to users through the notion of everywhere, anytime, and using any gadget. As a result, integrating networks is seen as the next big hurdle for digital communication to overcome [22]. Since conversations over LTE began inside 3GPP in 2004, it cannot be a very fresh field. An example of this "unceasing toil" is the hundreds of engineers who are involved with the 3GPP standards panels. Long-Term Evolution is the innovative technology that will utilize MIMO antenna transmission to





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push the boundaries of limit. The connection with the radio system and the construction of the radio infrastructure are both encompassed in LTE system layouts.

Media Independent changeover (MIH) is a continuing effort in the IEEE802.21 WG which involves many types of connections. Its goal is to create smooth changeover across numerous broadband networks, despite their equipment. [1, 2, 3, and 4]. Anyone can own the necessary LTE access network; what's important is that the right regulations and service level contracts be in place to ensure that users can roam and interact seamlessly with one another [5]. Because of its broad signal area, assistance with user independence, inherent quality of service, and high data rate capabilities, WiMax technology relying on the IEEE802.16e equipped has garnered a lot of interest from both consumers and broadband providers [34]. One possible way to achieve the 4G situation is by combining and interconnecting LTE and WiMax innovations [24].

The wireless connectivity Broadband Conformity Coalition, better known as WiFi [26], is responsible for verifying that devices that adhere to the IEEE 802.11 standard can communicate with one another. The maximum indoor visibility radius for a WiFi access point is around 100 meters [23]. The range of WiFi outside depends on factors like the weather and other external factors, although it may be as long as 300 meters [25]. From 11 Mbps to 54 Mbps, that's the range of WiFi data transmission speeds [3]. Known variously as Fixed WiMax and the IEEE802.16d protocol, it forms the basis of WiFi-WiMax integration. Additionally, the IEEE 802.16e regulation has not been implemented yet. In [30], the authors conduct a simulation research that demonstrates how connectivity impacts the efficiency of program technologies such as HTTP, FTP, email, and VoIP. In [31], we find a method for safely integrating wireless diverse networks. In [32], we find a standard infrastructure for adaptive radio-assisted integration of heterogeneous wireless networks. In reference [33], a method for vertical handoff in HetNet is described.

Simulation Environment

The objective of the investigation is to analyze the working of tight and loose coupling. And impact over the protocols of different layers. For application layer protocols Email, FTP, VOIP and http are considered. For individual network-based parameters WiMax and WLAN are considered. A hypothetical research with coupled WiMax/WLAN is conducted using the LTE Opnet version modeling. Twenty square kilometers constitute the space of the model. The experiment time is configured at 3600 seconds. The first graph shows the tightly coupled modeling situation. Simulation scenario for loose interface/ coupling is given in figure 2. Four application servers are connected with cloud and user from both scenarios for WiMax and WLAN are trying to download the files and services from the cloud. The users are mobile and speed is varying from 0-20m/s and following a random way point trajectory.

RESULT ANALYSIS AND DISCUSSION

Performance of Application Protocol

Implementation procedures provide the foundation for measuring the efficacy of networks for people who use it. Here we get into the topic in use protocols, specifically looking at Email, FTP, VOIP, and HTTP. The third figure through Figure 13 show the FTP's functionality. For the different kind of application tight coupling is showing better performance.

Performance of WiMax

Monitoring the efficiency for each person innovations and the effects of combining on these innovations are crucial during diverse system integration. See Figures 14 and 15 for WiMax effectiveness details. As a result of strong association, the WiMax layer duration is shown to be optimal. Keep in mind that the WiMax layer's strong productivity for informal coupling might be because of the regulation sessions.





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Performance of WLAN

You can see how well the WLAN infrastructure is doing in Figures 16 and 17. Though WLAN delay is good in case of loose coupling but it has been due to the internal performance but not for the integrated network. If we are integrating the network then we have to consider all possible scenarios and outcomes.

CONCLUSION

In this paper WiMax and WLAN are integrated under the umbrella of LTE. We simulated the results for Loose and Tight interfacing/coupling for designing the framework for the HetNets. And we observed that tight coupling is good in comparison to loose coupling. We can consider this point when laying the groundwork for technology integration. Further discussion may include the designing of Framework in order to integrate wireless networks that are not homogeneous.

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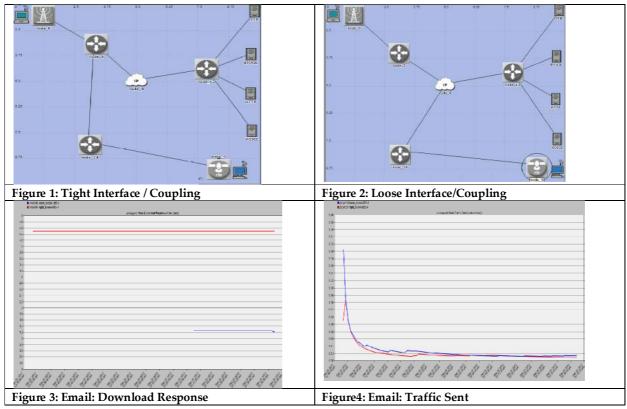




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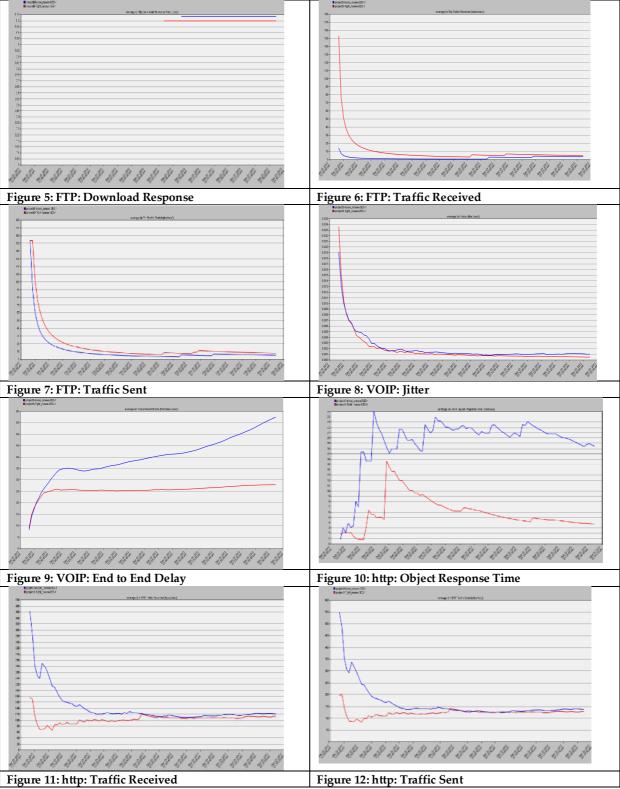
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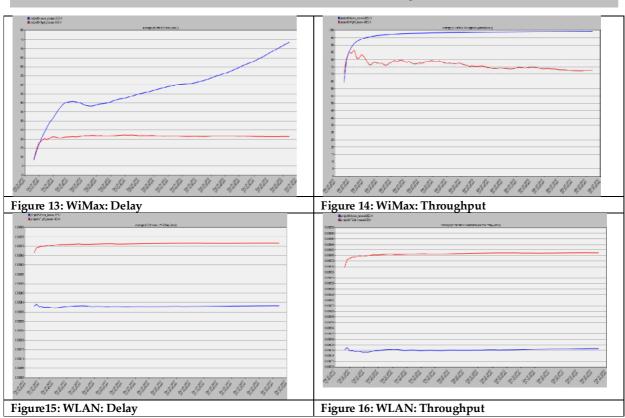
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RESEARCH ARTICLE

Descriptive Analytics On Healthcare Data -A Systematic Approach To **Enhance Patient Engagement**

Lavonya A1, Sunil Raj Y2, Charles S3 and A Leolin Arockiadass4

- ¹PG Student, Dept of Data Science, St. Xavier's College (Autonomous), Palayamkottai, India.
- ²Assistant Professor, Dept of Data Science, St. Xavier's College (Autonomous), Palayamkottai, India.
- ³Assistant Professor, Dept. of Computer Science, AL Musana Institute of Engineering & Technology, Sultanate of Oman, UAE
- ⁴Assistant Professor, Dept of Computer Science, St. Xavier's College (Autonomous), Palayamkottai, India.

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*Address for Correspondence Sunil Raj Y

Assistant Professor, Dept of Data Science, St. Xavier's College (Autonomous), Palayamkottai, India. Email: ysrsjccs@gmail.com



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ABSTRACT

Healthcare analytics leverages mathematical methodologies to analyze vast amounts of data, facilitating informed decision-making within the healthcare sector. By integrating data-driven insights, healthcare analytics enhances patient care outcomes and operational efficiencies across various healthcare settings. Its application spans from refining diagnostic processes to optimizing resource allocation and treatment strategies. The transformative impact of healthcare analytics parallels its counterparts in finance, marketing, and technology sectors, driving advancements in healthcare delivery. Despite its potential benefits, healthcare analytics faces challenges such as data privacy concerns and interoperability issues that necessitate innovative solutions. Addressing these challenges is essential to fully harness the potential of healthcare analytics in improving patient outcomes and organizational effectiveness.

Data Analytics, Descriptive Analytics, Diagnostic Analytics, Predictive Analytics, Prescriptive Analytics, Healthcare Analytics

INTRODUCTION

Data analytics plays a crucial role in extracting valuable insights from raw data, with its applications expanding significantly due to the widespread adoption of smart devices, which furnish abundant information about monitored





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assets. The temporal dimension of analytics empowers organizations to make informed decisions at specific junctures, profoundly impacting business performance metrics such as revenue, cost management, resource utilization, and operational efficiency. Integration of advanced analytical tools like machine learning (ML) and artificial intelligence (AI) further augments the efficacy of data analytics processes. Access to data at various managerial levels facilitates streamlined monitoring of processes. Depending on the timing of execution, data analytics finds diverse applications, broadly categorized as descriptive, diagnostic, predictive, and prescriptive analytics. Healthcare represents society's collective endeavor to ensure, finance, provide, and promote health, with a paradigm shift in the twentieth century towards wellness and preventive measures. Standardized protocols within healthcare aid in evaluating actions or situations influencing decision-making processes. As a multi-dimensional system, healthcare aims to diagnose and treat illnesses or disabilities, emphasizing the fundamental goal of fostering overall well-being. The following sections are organized as; background of the study, review of literature, analysis on the existing work, issues discussed and amble solutions for the issues and lastly the conclusion.

Background

Healthcare analytics, a specialized branch of data analytics, harnesses historical and current data to extract actionable insights, refine decision-making processes, and optimize outcomes within the healthcare domain. Beyond merely benefiting healthcare organizations, this analytical approach plays a pivotal role in enhancing the overall patient experience and advancing health outcomes. Given the vast reservoir of detailed records pervasive throughout the healthcare industry, a wealth of valuable data is readily available, presenting abundant opportunities for thorough analysis and interpretation.

Figure 1, presents a visual representation of how healthcare analytics fits within the field of data analytics. It illustrates how healthcare analytics is a specialized subset of data analytics, focusing specifically on the healthcare sector. Healthcare analytics, including descriptive, diagnostic, predictive, and prescriptive analytics, play a vital role in improving patient care for those with End Stage Renal Disease (ESRD). Descriptive analytics helps understand past trends, while diagnostic analytics identifies root causes of specific events. Descriptive analytics provides insights into historical events, addressing queries like the number of hospitalized patients or the percentage discontinuing home therapy. Diagnostic analytics delves into understanding causal factors behind events, addressing queries like why patients visited hospitals, abandoned home therapy, or failed to meet treatment targets [11]. Predictive analytics is a powerful tool in healthcare, leveraging historical data to forecast future events with notable accuracy. These can predict patient transitions, such as those likely to shift from home therapy to in-center care, facilitating resource allocation and patient support planning [12]. Healthcare analytics contributes significantly to improving patient outcomes by identifying trends and patterns for early disease detection and timely intervention, ultimately enhancing patient well-being. Personalized medicine is facilitated through the analysis of patient data, allowing for the development of tailored treatment plans that consider individual health characteristics, thus improving the efficacy of medical interventions and patient satisfaction [13]. Additionally, analytics plays a crucial role in fraud detection, ensuring fairness and accuracy in financial transactions, which is essential for maintaining trust and integrity within the healthcare system. Healthcare analytics significantly enhances the quality of patient services by providing insights into industry trends and practices. Descriptive analytics, known for its ability to unveil prevailing facts, is particularly suited for examining healthcare data.

REVIEW OF LITERATURE

Descriptive analytics involves the exploration and interpretation of historical data to identify patterns and trends. In the healthcare context, it encompasses various components such as data visualization, summarization, and reporting techniques. Studies by Jeremiah O A, et al. (2024) highlight the importance of understanding the foundational aspects of analytics in healthcare settings[14]. Numerous studies have demonstrated the diverse applications of descriptive analytics in healthcare. Research by F AI Zoubi, et al. showcases how descriptive analytics can be employed to monitor patient health, optimize resource allocation, and improve overall operational efficiency within healthcare





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organizations [15]. The dynamic nature of healthcare demands a constant evolution of analytic approaches. Research by Lee and Kim., 2023, explores emerging trends such as the integration of artificial intelligence and machine learning techniques with descriptive analytics, offering insights into potential advancements in the field. Sabah M, et al, 2022, have focused on Health data analytics and identified the improvements it makes towards patient care, faster diagnoses, preventive measures, more personalized treatment and provide more informed decision-making. Also, it was found that healthcare analytics could assist improving patient outcomes, enhancing care management, and address social determinants of health [7, 11]. As the health analytics grows eventually in the healthcare space, healthcare organizations could gather data from manifold sources, after which apply data analytics, and provide insights for the population.

METHODOLOGY

Healthcare analytics has emerged as a pivotal tool in improving patient care and operational efficiency within the healthcare domain. This proposed methodology outlines a structured approach to conducting research in healthcare analytics, aiming to address existing research issues and provide actionable solutions. The methodology integrates various analytical techniques and data-driven approaches to inform decision-making processes in healthcare settings. The first step in our proposed methodology is to identify and define the specific healthcare problem or challenge to be addressed through analytics. This may include issues such as optimizing resource allocation, improving patient outcomes, reducing costs, or enhancing operational efficiency. Once the problem is identified, relevant data sources need to be identified and collected. This may include electronic health records (EHRs), medical imaging data, administrative data, patient surveys, and other relevant sources. Data cleaning and preprocessing techniques are then applied to ensure data quality and consistency. EDA involves analyzing and visualizing the collected data to gain insights into underlying patterns, trends, and relationships. Descriptive statistics, data visualization techniques, and exploratory data mining methods are employed to identify potential correlations and associations within the data. Predictive modeling techniques, such as machine learning algorithms, are utilized to build predictive models based on the available data. These models can be used to forecast future healthcare outcomes, predict patient risk levels, or identify potential areas for intervention. Figure 2, describe the methodology used to analyze the patient engagement scenario for reducing the complication ratio and reducing the unnecessary visits. After the exploration of data, models are developed after which it is evaluated and validated using appropriate metrics and validation techniques. This ensures the reliability and generalizability of the models and their applicability to real-world healthcare scenarios. The insights gained from the analytics process are interpreted in the context of the healthcare problem being addressed. Key findings and actionable insights are identified, which can inform decision-making processes and drive improvements in patient care and healthcare operations. The validated models and insights are then implemented into healthcare practice, either through integration into existing systems or through the development of new tools and workflows. Close collaboration with healthcare stakeholders is essential to ensure successful implementation and adoption. Finally, the implemented solutions are continuously monitored and evaluated to assess their impact on healthcare outcomes and operations. Iterative refinements and improvements are made based on feedback and new data, ensuring ongoing optimization and effectiveness.

RESULTS AND DISCUSSION

The dataset chosen consists features of about 300k, which records the medical appointments of patients. The focus is to analyze whether the patients showed up for their appointments and the factors behind their response. The dataset consists of 15features, including appointment details, medical conditions, and appointment attendance status. Age and gender represent demographic information, while Appointment Registration and Appointment Date indicate scheduling details. The Status column distinguishes between patients who attended their appointments and those who missed them. Medical conditions such as diabetes, alcoholism, hypertension, handicap, smokers, tuberculosis, and whether the patient has a medical scholarship are also recorded. SMS_ Reminder indicates if the patient received reminder messages. Awaited Time represents the duration patients had to wait from registration to appointment.





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This data allows for analysis of factors influencing appointment attendance and the effectiveness of SMS reminders. Figure 3 displays the count of patient show-ups categorized by various factors such as age, gender, medical conditions (diabetes, alcoholism, hypertension, handicap, smokers, tuberculosis), presence of medical scholarship, receipt of SMS reminders, and waiting time. This visualization provides insights into which demographics or conditions correlate with higher or lower attendance rates, aiding in the identification of factors influencing appointment attendance. Such analysis can inform targeted interventions to improve attendance and patient outcomes.

Figure 4 presents a gender distribution plot that depicts the count of appointments by gender, further segmented by appointment status (attended or missed). This visualization offers a comprehensive view of appointment attendance patterns across different genders. By visualizing the distribution of show-ups and no-shows within each gender category, healthcare providers can gain insights into potential gender-specific factors influencing appointment adherence. This understanding facilitates the development of targeted interventions to address any disparities and improve overall appointment attendance rates. Figure 5 displays a count plot based on the "Day Of The Week" variable, illustrating the distribution of appointments across different days of the week. This visualization provides insights into appointment scheduling patterns, revealing which days are most popular for appointments and which days have lower appointment volumes. Analyzing this information enables healthcare providers to optimize appointment scheduling and resource allocation, ensuring efficient and effective healthcare service delivery throughout the week.

Figure 6 depicts the percentage of missing values across different variables in the dataset. This visualization offers insights into the completeness of the data and highlights any variables with significant missing values. Understanding the extent of missing data is crucial for ensuring the reliability and validity of analytical results. By identifying variables with high percentages of missing values, researchers can prioritize data collection efforts or employ appropriate imputation techniques to mitigate the impact of missing data on subsequent analyses. Figure 7 illustrates the status of patients who turned up for their appointments, with each status. Also, the visualization provides a nuanced understanding of the various factors and circumstances influencing appointment attendance, highlighting the diverse pathways through which patients ultimately turn up for their scheduled appointments.

Comparison and Analysis

The descriptive analysis is conducted on the chosen patient data set where it is free from missing values. Among the chosen set of data about 69.75% of people showed-up and the remaining 30.24%did not show up for their appointments. The analysis shows that female patients have taken more appointments than male patients. There are around 200505 Females and out of them around 70.13% showed-up for their appointments and the remaining 29.87% did not show up. Out of 99495 males, 69% showed-up while 31% did not show up. The highest no of people showed-up was between the age group 41-61 and between the age group between 101-121 people did not show up for their appointments. The least number of appointments were made on Sunday while most of them are on Wednesday The observed results in figure 8, shows that the proposed methodology exhibits better based on the size of data chosen. As the impact the complication ratio is comparatively reduced to 98% while the existing proposal showed 92%. Also, the number of unnecessary visits of the patients is also reduced and it was up to 89%. This shows the importance of analytics in healthcare as it substantiates the assistants by enhancing the overall performance, while provide better service.

CONCLUSION

Healthcare analytics stands as a transformative tool in enhancing patient care and operational efficiency within the healthcare sector. The proposed methodology delineates a systematic approach to healthcare analytics research, aimed at tackling prevailing research challenges and furnishing actionable solutions. By amalgamating diverse analytical techniques and data-driven methodologies, informed decision-making in healthcare settings becomes





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attainable. Initial findings from patient data analysis reveal that about 69.75% of individuals attended their appointments while 30.24% did not, underscoring the significance of attendance trends. Gender-based analysis demonstrates that female patients made more appointments (200,505) compared to males (99,495), with 70.13% of females showing up versus 69% of males. Age-group specific analysis indicates the highest attendance within the 41-61 age range, while those aged 101-121 exhibited lower attendance rates. Additionally, appointment distribution analysis highlights a preference for midweek appointments, with fewer appointments scheduled for Sundays. Therefore, it is evident that insights from the descriptive analytics drives informed decision-making. The implementation of validated solutions in healthcare, enable continuous refinement to optimize scheduling, and service delivery for improved patient outcomes and operational excellence.

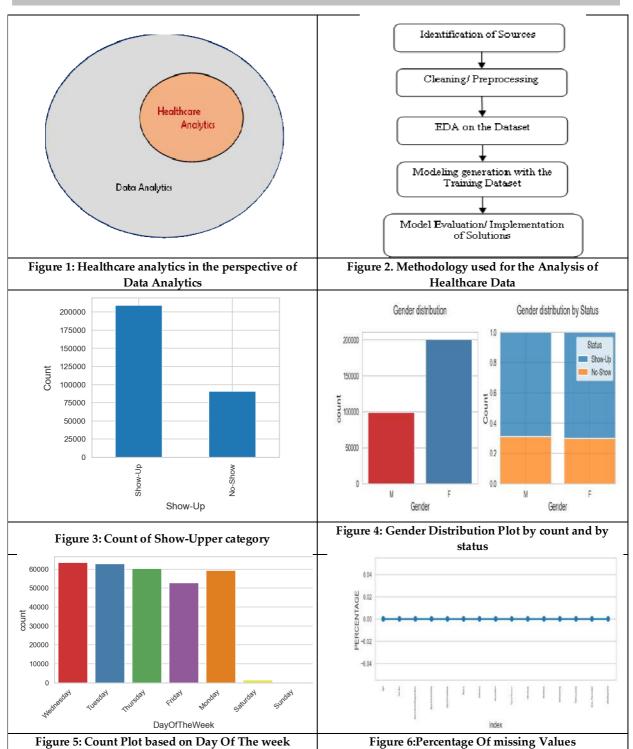
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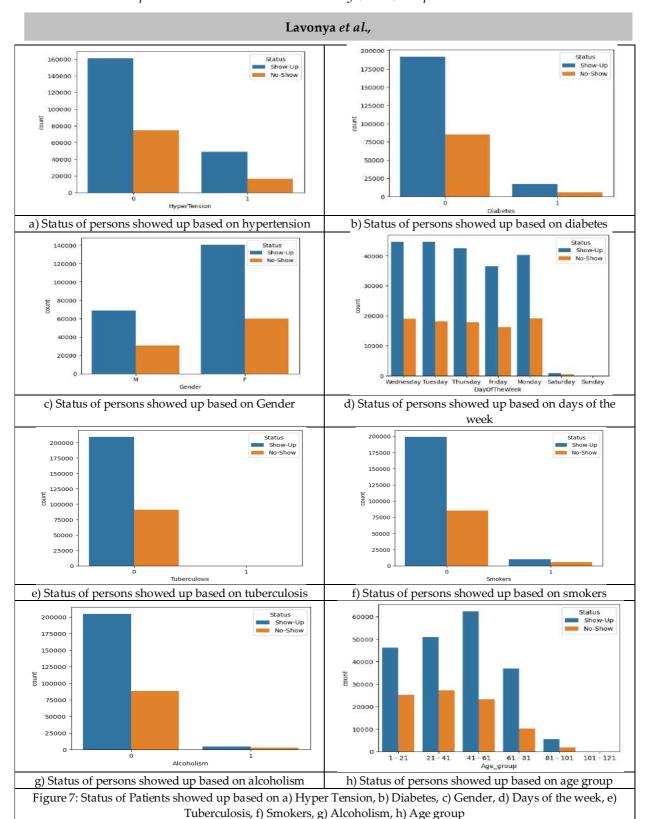


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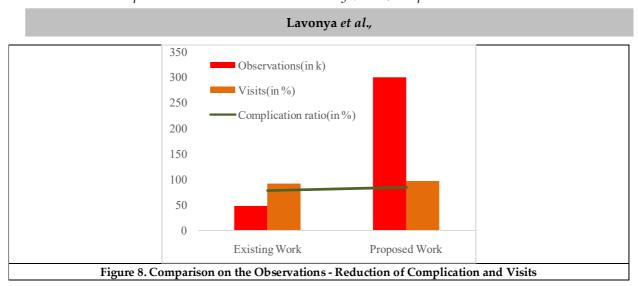
















RESEARCH ARTICLE

BER Performance Evaluation of DWDM Optical Network at Various **Channel Spacing**

Poorna Lakshmi Uppalapati1*, Kotika Sai Kumar2, Lingaiahgari Amulya2 and Arige Sai Yashwanth²

¹Professor and HoD, Department of Electronics and Communication Engineering, Vignana Bharathi Institute of Technology, Hyderabad, Telangana, India.

2Student, Department of Electronics and Communication Engineering, Vignana Bharathi Institute of Technology, Hyderabad, Telangana, India.

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*Address for Correspondence Poorna Lakshmi Uppalapati

Professor and HoD,

Department of Electronics and Communication Engineering,

Vignana Bharathi Institute of Technology,

Hyderabad, Telangana, India

Email: uppalapati.poornalakshmi@vbithyd.ac.in



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ABSTRACT

Dense Wavelength Division Multiplexing (DWDM) is a method of optical fiber transmission that employs densely spaced light frequencies and can simultaneously transport large amounts of data. This is a useful method to increase the system's capacity. In this work, we studied the 16 x 10 GBps DWDM communication network over a 50 km length Single-Mode Fiber (SMF) at various channel spacing, including 0.8 nm and 1.6 nm with bit rates of 10 Gbps.

Keywords: DWDM, Single-Mode Fiber, Dispersion-Compensating Fiber, Channel Spacing

INTRODUCTION

The development of optical communication technologies has been accelerated by the expansion of data-hungry applications and the unrelenting demand for higher network capacity [1]. Dense Wavelength Division Multiplexing (DWDM) has become a key option, enabling the simultaneous transmission of numerous data channels over a single optical fiber while utilizing various light wavelengths [2]. The performance of the DWDM optical network can fluctuate significantly when the channel spacing, which is the frequency difference between neighboring channels, varies [3, 4]. In This work, the Bit Error Rate (BER) of a DWDM optical network over a range of channel spacing values [5] is studied. A key indicator for evaluating the DWDM optical network is BER. It measures the proportion of wrongly received bits to all transmitted bits, demonstrating the robustness of the system against errors and the overall data integrity. It is possible to learn a lot about how the network reacts to changes in the spectral allocation of channels by methodically





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analyzing the BER performance at various channel spacing configurations[6]. The study entails a thorough examination through simulation. The effect of various channel spacing values on the BER performance are investigated, revealing information on the network's capability to maintain dependable data transmission under various circumstances [7, 8, 9]. The results of the study are expected to aid in the optimization of DWDM optical networks and in the development of more reliable and effective communication systems that can handle the increasing demands of contemporary datacentric environments[10, 11].

Schematic of DWDM system

Fig. 1 shows DWDM communication system. Multiple data at the output of transmitters is given to multiplexer, which is used to combine multiple data and transmit over an optical fibre. The signal is then given to an EDFA (erbium-doped fibre amplifier). The optical demultiplexer output is given to respective receivers.

BER Performance Analysis

BER performance is evaluated for channel spacing of 0.8 nm and 1.6 nm for 2, 4, 8 and 16-channel DWDM optical network. PRBS generator generates pseudo random bit sequence used to modulate the light carrier in the transmitter section. The modulated carrier signals are multiplexed using Multiplexer and placed on an optical fiber channel. At the other end of the optical fiber the wavelength selection filter selects and receiver detects the electrical signal. Power normalizer is used to normalize the power levels. BER tester is used to analyse the Bit error rate. Here, we're using two channels operating at a data rate of 10 Gbps. RSoft Otsim is uesd for the simulation of DWDM network. Table 1 shows BER values for 2 channel.

DWDM with channel spacing 0.8 nm and Table 2 shows BER values for 2 channel DWDM with channel spacing 1.6 nm. Fig. 3 shows log(BER) value for two channel DWDM system with channel spacing 0.8 nm and 1.6 nm. BER values for DWDM system with 1.6 nm channel spacing are better than 0.8 nm channel spacing. The same can be seen for 4, 8 and 16 channel DWDM system.

Table 3 shows BER values for 4 channel DWDM with channel spacing 0.8 nm and Table 4 shows BER values for 4 channel DWDM with channel spacing 1.6 nm. Fig. 4 shows log (BER) value for four channel DWDM system with channel spacing 0.8 nm and 1.6 nm. Similarly, the BER performance is evaluated for channel spacing 0.8 nm and 1.6 nm for 8 channel and 16 channel DWDM. Table ?? shows BER values for 8 channel DWDM with channel spacing 0.8 nm and Table ?? shows BER values for 8 channel DWDM with channel spacing 1.6 nm. Fig. 5 shows log (BER) value for eight channel DWDM system with channel spacing 0.8 nm and 1.6 nm. Table 7 shows BER values for 16 channel DWDM with channel spacing 0.8 nm and Table 8 shows BER values for 16 channel DWDM with channel spacing 1.6 nm. Fig. 6 shows log (BER) value for sixteen channel DWDM system with channel spacing 0.8 nm and 1.6 nm. It can be observed that Bit Error Rate decreases as channel spacing is increased. At the chosen parameters, data rate 10 Gbps, NRZ modulation and link length 50 km the BER values are far better than threshold value 10–12. For a DWDM network with channel spacing 1.6 nm, the BER value is even better than 0.8 nm channel spacing so the data rate can be increased.

CONCLUSION

Through this simulation, how different channel spacings influence the performance of a 2,4,8,16 channel DWDM system has been studied. BER performance evaluated at channel spacing 0.8 nm and 1.6 nm. BER value is better for channel spacing 1.6 nm for all the 2,4,8 and 16 DWDM networks. The BER value for the chosen simulation parameters is far better than the trshold value of BER 10–12. Therefore the data rate can be allowed to increase. Further the DWDM network with more number of channels, channel spacing and data rate can be studied.





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Table 1: 2-channel with spacing of 0.8.

Wavelength	BER Value	
1552.52	7.68e-13	
1553.32	8 6.87e-20	

Table 2: 2-channel with spacing of 1.6.

Wavelength	BER Value	
1552.52	6.13e-25	
1554.12	6.87e-29	

Table 3: 4-channel with spacing of 0.8.

Wavelength	BER Value	
1552.52	9.37e-15	
1553.32	1.13e-11	
1554.12	4.15e-10	
1554.92	8.12e-20	





Table 4: 4-channel with spacing of 1.6.

Wavelength	BER Value	
1552.52	7.50e-15	
1554.12	7.65e-27	
1555.72	6.23e-20	
1557.32	1.4764e-32	

Table 5: 8-channel with spacing of 0.8.

Wavelength	BER Value	
1552.52	2.64e-15	
1553.32	1.54e-12	
1554.12	8.66e-11	
1554.92	2.19e-12	
1555.72	1.56e-11	
1556.52	1.05e-10	
1557.32	2.05e-10	
1558.12	8.21e-18	

Table 6: 8-channel with spacing of 1.6.

Wavelength	BER Value	
1552.52	2.64e-15	
1553.32	1.54e-12	
1554.12	8.66e-11	
1554.92	2.19e-12	
1555.72	1.56e-11	
1556.52	1.05e-10	
1557.32	2.05e-10	
1558.12	8.21e-18	





Table 7: 16-channel with spacing of 0.8.

Wavelength	BER Value
1552.52	1.81e-15
1553.32	3.8e-13
1554.12	1.24e-10
1554.92	1.42e-20
1555.72	4.47e-12
1556.52	3.16e-11
1557.32	6.57e-11
1558.12	6.24e-12
1558.92	4.92e-11
1559.72	4.03e-12
1560.52	6.56e-12
1561.32	3.13e-10
1562.12	3.38e-11
1562.92	3.09e-09
1563.72	1.03e-10
1564.52	1.04e-14

Table 8: 16-channel with spacing of 1.6.

Wavelength	BER Value
1552.52	9.35e-22
1554.12	9.94e-27
1555.72	5.05e-20
1557.32	1.38e-29
1558.92	3.67e-24
1560.52	5.62e-21
1562.12	1.45e-22
1563.72	2.36e-25
1565.32	6.75e-20
1566.92	1.00e-25
1568.52	5.79e-21
1570.12	1.53e-16
1571.72	3.49e-19
1573.32	4.68e-15
1574.92	8.04e-18
1576.52	4.28e-20





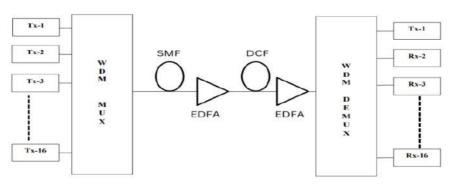


Figure 1: Schematic Diagram of DWDM System

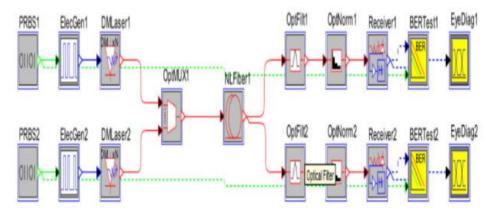


Figure 2: Simulation set up for 2-channel DWDM Optical fiber link

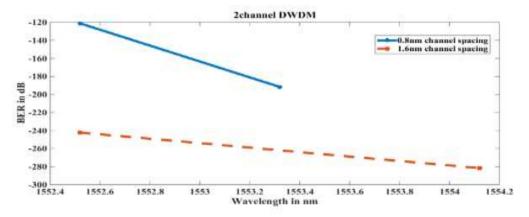


Figure 3: BER values for two channel DWDM with 0.8 and 1.6 spacing





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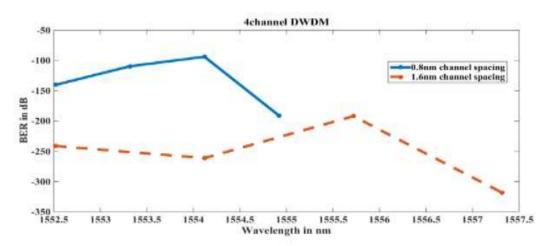


Figure 4: 4-channel Performance evaluation of BER between 0.8 and 1.6 spacing

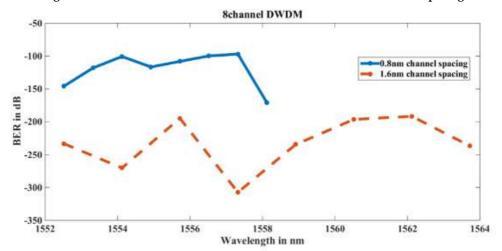


Figure 5: 8-channel Performance evaluation of BER between 0.8 and 1.6 spacing

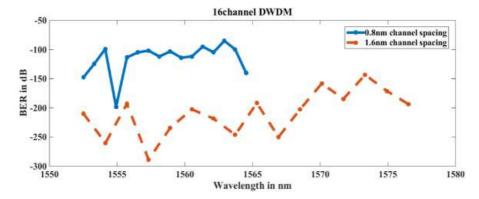


Figure 6: 16-channel Performance evaluation of BER with 0.8 and 1.6 nm channel spacing





RESEARCH ARTICLE

Taxonomic Enumeration of Natural Dyes from India: Sir Thomas Wardle's Collection

Sudeshna Datta¹, Surendra Kumr Sharma², Manas Bhaumik³ and Kangkan Pagag^{4*}

¹Botanist, Botanical Survey of India, Central National Herbarium, Howrah, West Bengal, India.

²Botanical Assistant, Botanical Survey of India, Central Botanical Laboratory, Howrah, West Bengal, India.

³Scientist-F, Botanical Survey of India, Industrial Section Indian Museum, Kolkata, West Bengal, India.

⁴Assistant Professor, Department of Botany, Sibsagar College, Joysagar (Autonomous), (Affiliated to Dibrugarh University), Assam, India.

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*Address for Correspondence

Kangkan Pagag

Assistant Professor,

Department of Botany, Sibsagar College, Joysagar (Autonomous),

(Affiliated to Dibrugarh University),

Assam, India.

Email: kpagag@gmail.com



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ABSTRACT

The present study deals with archival data of Indian plants used by renowned British dyer & innovator Sir Thomas Wardle. He had used Indian plants singly or in combination which brought out various shades of colors on various fabrics viz. cotton, silk, wool. He had made sets of catalogues of samples dyed with natural dyes and one is deposited at Industrial Section Indian Museum, Kolkata. Therefore the present systematic taxonomic enumeration will help dyers or weavers or handloom industry in knowing dye yielding plants used in 19th century in India which may promote sustainable organic dye.

Keywords: Economic, colors, fabric, organic, sustainable.

INTRODUCTION

Sir Thomas Wardle (1831-1909), a remarkable British dyer and printer, innovator and entrepreneur who was erudite in dyeing of silk textiles and bringing out new innovations for it. He had a deep passion for silk and dyes. And due to this passion he was dedicated towards wild silks and dyes particularly Indian which created a platform for Indian dyes and silks in India & Europe. He was also a president of the Silk Association of Great Britain and contributed in setting up a silk factory in 1897. Sir Thomas Wardle studied wild silks of India (1880). He was reputed in bringing out different shades of color to textiles. He received a knighthood and many awards due to his marvelous feat of





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researches. He was also granted a Fellowship of the Royal Society of Chemists. India at 19th century was rich in textile industry and many of the products or textiles were plants based. Indian Indigo was immensely popular in Europeans. Europe during those period was exported Indian textiles with Indian dyes. Thomas Wardle was drawn towards the richness of the natural dye yielding plants. He had experimented with many Indian dyes. In 1878 he had started the exploring the potentials of tinctorial powers and properties many of Indian dyes on the request of Government of India. The then Secretary of State for India had stated to him that if the natives of India could be taught how to dye their Tusser silk, a valuable industry could be created for their benefit (Wardle, 1887). These dyes were plant based, minerals as well as insects. To carry out such a huge task was very tedious and time consuming and it took many years to bring out the samples and present the report. He was helped by Mr Hugh Mccain who visited or brought out the information from the common locals or dyers also by providing raw materials of Indian dyes. Mr Hugh Mccain (1883) had published these details on Dyes of Bengal too. But he had just published the data he obtained from the field. It was Sir Wardle who made the album of the samples of the dyed textiles out of his experimentation. He chose different fabrics such as cotton, wool, muslin, silk, tussar for dyeing. He employed all the known methods of dyeing. He had used not only known mordants but used some of his private methods or mordants and as a result he subsequently discovered others. Since he was in business of dye and print he didn't reveal the methodology or percentage of mordant and dyes used. He used fustic also. He deposited the catalogue to the Government and published the report (1887). These albums of samples which were in possession of Industrial Section Indian Museum still attracts many researchers and dye experts from all over the globe. The samples are still vibrant in colour and which shows the efficacy of Indian dyes as well as working techniques of Wardle. According to Wardle some of the Indian dyes possessed distinct properties and in many cases superior to the dyes obtained from artificial sources. The set of 3000 samples are consists of fifteen volumes and arranged 09 samples per page with names of the dye and mordant used with common or scientific name of the plant. One will be mesmerized when goes through these samples which are the beautiful shades of pink, red, yellow, green, blue, brown etc. Therefore it reflects the importance of Indian dyes along with the techniques used in dyeing.

MATERIALS AND METHODS

The catalogues more than century old & prepared by Sir Thomas Wardle possessed by Industrial Section Indian Museum, Botanical Survey of India were consulted. Also the report submitted by Thomas Wardle (1887) to Government of India consulted & literature by M'cann (1883), were studied. Photographs taken. Enumeration of the plants was prepared with the current scientific nomenclature by consulting POWO (2023).

RESULT

Along with the plant dye, the role of mordants are very important in production of particular shades. Wardle had only written coded symbols for mordants which is yet to be deciphered. The plants used by him were the dye yielding plants of that period which included many common plants. The following is the enumeration of dye-yielding plants along with the scientific name, parts used and colour shades

Sl. No.	Common Name	Scientific Name	Parts Used	Colour shades obtained
1	Indigo	Indigofera tinctoria L.	Whole plant	Green, Blue, Blue- green
2	Madder	Rubia cordifolia L.	Bark, Root	Red, Brown, Maroon, Violet, Orange, Peach, Pink, Purple, Grey
3	Weld	Reseda luteola L.	Whole plant	Yellow, Grey, Green





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4	Sapan wood	Caesalpinia sappan L.	Wood	Coral, Brown, Peach, Yellow, Red, Maroon, Orange
5	Sanders wood	Pterocarpus santalinus L. f.	Wood	Brown, Red
6	Annatto	Bixa Orellana L.	Seed	Yellow
7	Turmeric	Curcuma longa L.	Rhizome	Brown, Blue, Yellow, Off-white
8	Safflower	Carthamus tinctorius L.	Flower	Brown, Orange, Pink, Yellow, Red
9	Manjeeth	Rubia manjith Roxb.	Root	Brown, Yellow, Orange, Red, Maroon, Green, Magenta, Peach
10	Berberry	Berberis lyceum Royle & Berberis tinctoria Lesch.	Wood, Root	Yellow, Brown, Yellow-ochre, Off- white, Khaki
11	Peepal	Ficus religiosa L.	Bark	Ochre, Brown, Golden yellow, Yellow, Off- white
12	Pulha Babul, Kekir, Babool	Vachellia nilotica subsp. tomentosa (Benth.) Kyal & Boatwr. (syn. Acacia Arabica)	Bark, Pods	Brown, Yellow-ochre
13	Lodh	Symplocos racemosa Roxb.	Bark	Yellow-ochre, Brown, Off-white
14	Jigna	Lannea coromandelica (Houtt.) Merr. (syn. Odina wodier)	Bark	Brown
15	Himalayan Blue Pine	Pinus wallichiana A.B. Jacks (Pinus excelsa)	Bark	Brown
16	Chir pine	Pinus roxburghii Sarg. (Syn. Pinus longifolia)	Bark	Brown
17	Jack wood	Cryptocarya glaucescens R. Br.	Wood	Brown
18	Peach Tree	Prunus persica (L.) Batsch	Root bark	Brown
19	Daruharidra, Al roots, Bhartondi, Meddi tree	Morinda citrifolia L.	Root, Bark	Yellow, Brown, Off- white, ochre, Orange
20	Henna	Lawsonia inermis L.	Leaves	Yellow, Brown
21	Jangali badam	Terminalia catappa L.	Bark, Leaves	Yellow, Brown, Off- white, Yellow-ochre
22	Taira	Libidibia coriaria (Jacq.) Schltdl. (Syn. Caesalpinia coriaria)	Pods	Brown, Yellow-Brown
23	Chaikatha	Piper retrofractum Vahl (Syn. Piper chaba)	Not known	Brown, Yellow, Off- white
24	Ashna	Terminalia tomentosa Wight & Arn.	Bark	Brown
25	Tarwad, Tarota	Senna auriculata (L.) Roxb. (Syn. Cassia auriculata)	Bark, Leaves,Seeds	Brown, Yellow
26	Bokal	Mimusops elengi L.	Bark	Brown, Off-white
27	Madar	Erythrina Americana Mill. (Syn. Erythrina fulgens)	Bark	Brown, Yellow
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28	Bihul	Grewia optiva J. R. Drumm. ex Burret (Syn. Grewia oppositifolia)	Leaves	Off-white, Brown
29	Jangli erandi	Jatropha glandulifera Roxb.	Leaves	Yellow, Brown, Off- white
30	Anjan	Memecylon coeruleum Jack (Syn. Memecylon tinctorium)	Leaves	Yellow, Brown
31	Gab	Diospyros malabarica (Desr.) Kostel.	Leaves, Fruits	Yellow, Off-white
32	Bhauri	Symplocos theifolia D. Don	Leaves	Off-white, Khaki, Golden, Yellow
33	Arjun	Terminalia arjuna (Roxb. ex DC.) Wight & Arn.	Bark	Yellow, Brown
34	Pomergranate rind	Punica granatum L.	Rind	Brown, Yellow-ochre, Steel gray, Blue-Black, Yellow
35	Anglithor	Euphorbia tirucalli L.	Not known	Yellow, Brown, Off- white, Green
36	Harsinghar	Nyctanthes arbor-tristis L.	Flowers	Yellow
37	Toon	Cedrela toona Roxb. ex Rottler & Willd. (Syn. Cedrela toona)	Flowers, Bark	Yellow, Green, Brown, Off-white, ochre
38	Palas	Butea monosperma (Lam.) Kuntze (Syn. Butea frondosa)	Flowers	Yellow-ochre, Brown, Yellow, off-white, Ochre, Red
39	Paras pipal	Thespesia populnea (L.) Sol. Ex Corrêa	Flowers, Dried capsules and calyces	Yellow-ochre, Yellow
40	Dhao ka phul, Dhauri	Woodfordia fruticosa (L.) Kurz (Syn. Grislea tomentosa)	Flowers, Leaves	Yellow-ochre, Red, Yellow
41	Mittikut	Senegalia catechu (L.f.) P.J. H Hunter & Mabb. (Syn. Acacia catechu)	Catechu	Brown, Yellow, Blue- brown, Brown-red, Blue, off-white
42	Mahuwa	Madhuca longifolia var. latifolia (Roxb.) A. Chev.	Bark	Brown
43	Mango	Mangifera indica L.	Bark, Fruit pulp	Brown, Yellow, Yellow-ochre, Off- white
44	Kachnar	Bauhinia variegata L.	Bark	Brown, Yellow-ochre, Moss green, Off-white
45	Harra	Terminalia chebula Retz.	Bark, Nut	Yellow-ochre, Yellow, Brown
46	Amla	Phyllanthus emblica L.	Fruits, Leaves	Brown, Yellow-ochre
47	Mountain ru	Casuarina junghuhniana Miq. (Syn. Casuarina muricata)	Bark	Yellow-ochre, Yellow, Grey
48	Genda	Tagetes erecta L.	Flowers	Brown, Yellow
49	Amaltas	Cassia fistula L.	Bark	Brown, Biscotti
50	Dhaura	Terminalia anogeissiana Gere & Boatwr.	Leaves	Brown





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51	Rohun	Soymida febrifuga (Roxb.) A. Juss.	Bark	Yellow, Brown, Green, Off-white
52	Logwood	Haematoxylum campechianum L.	Extract	Violet, Blue, Brown, Grey
53	Barwood	Baphia nitida G. Lodd.	Notknown	Yellow, Brown, Maroon, Red, Violet, Pink
54	Fustic	Maclura tinctoria (L.) D. Don ex G. Don	Extract	Brown, Yellow, Green
55	Kamela	Mallotus philippensis (Lam.)Müll.Arg.	Fruits	Yellow
56	Weniwel	Coscinium fenestratum (Gaertn.) Colebr.	Wood	Yellow
57	Rang	Dicliptera tinctoria (Nees) Kostel (Syn. Peristrophe tinctoria)		Yellow, Brown, Peach, Pink
58	Red Creeper	Ventilago madraspatana Gaertn.	Root bark with galls	Brown
59	Tartigyan	Senna torra (L.) Roxb.	Seeds	Brown
60	Pista	Pistacia vera L.	Flowers	Brown
61	Dhayati, Kapila	Woodfordia fruticosa (L.) Kurz.	Flowers	Yellow
62	Suranji	Mammea suriga (BuchHam ex Roxb.) Kosterm (Syn. Calysaccion longifolium)	Root bark	Yellow
63	s.n.	Amaranthus tricolor L. (Syn. Amaranthus polygamous)	Stems	Yellow
64	Guljullil	Delphinium saniculifolium Boss.		Yellow
65	Cotton	Gossypium herbaceum L.	Flowers	Yellow, Brown
	Som	e combinations of plants for extraction of	dyes by Wardle	
66	Indigo and Weld	Indigofera tinctoria L., Reseda luteola L.	Whole plant	Prussian blue, Green, Blue
67	Indigo and Turmeric	Indigofera tinctoria L., Curcuma longa L.	Whole plant, Rhizomes	Yellow, Green, Blue- green, Blue.
68	Indigo and Berberry Wood	Indigofera tinctoria L., Berberis lyceum Royle/Berberis tinctoria Lesch.	Whole plant, Root	Green
69	Indigo and Logwood	Indigofera tinctoria L., Haematoxylum campechianum L.	Whole plant, Extract	Green, Blue-green
70	Indigo, Madder and Turmeric	Indigofera tinctoria L., Rubia cordifolia, Curcuma longa L.	Whole plant, Root, Rhizomes	Brown, Green
71	Indigo, Berberry Wood and Munjeet	Indigofera tinctoria L., Berberis lyceum Royle/Berberis tinctoria Lesch., Rubia munjith Roxb. ex Fleming	Whole plant, Root	Blue, Green
72	Indigo, Madder and Berberry Wood	Indigofera tinctoria L., Rubia tinctorum L., Berberis lyceum Royle/Berberis tinctoria Lesch.	Whole plant, Root	Green
73	Indigo, Madder and Sanderswood	Indigofera tinctoria L., Rubia cordifolia L., Pterocarpus santalinus L. f.	Whole plant, Root, Wood	Blue
74	Indigo, Logwood and Berberry Wood	Indigofera tinctoria L., Haematoxylum campechianum L., Berberis lyceum Royle	Whole plant, Extract, Root	Green





75	Indigo, Madder, Sanderswood and Turmeric	Indigofera tinctoria L., Rubia cordifolia L., Pterocarpus santalinus L.f., Curcuma longa L.,	Whole plant, Root, Wood, Rhizomes	Brown, Yellow, Green
76	Indigo, Berberry Wood, Turmeric and Munjeet	Indigofera tinctoria L., Berberis lyceum Royle/Berberis tinctoria Lesch., Curcuma longa L., Rubia munjith Roxb. ex Fleming	Whole plant, Root, Rhizomes, Root	Green
77	Indigo, Fustic, Madder and Barwood	Indigofera tinctoria L., Maclura tinctoria (L.) D. Don ex Steud., Rubia cordifolia L., Baphia nitida Lodd.	Whole plant, Extract, Root	Green, blue, olive green
78	Indigo, Berberry Wood, Logwood and Madder	Indigofera tinctoria, Berberis lycium, Haematoxylum campechianum, Rubia cordifolia L.	Whole plant, Root, extract, Root	Brown
79	Manjeeth and Palas	Rubia munjith Roxb. ex Fleming, Butea monosperma (Lam.) Kuntze	Root, Flowers	Brown, Yellow
80	Madder and Turmeric	Rubia cordifolia L., Curcuma longa L.	Root, Rhizomes	Brown
81	Madder and Sanderswood	Rubia cordifolia L., Pterocarpus santalinus L. f., Butea monosperma (Lam.) Kuntze	Root, Wood	Brown, Maroon, Orange
82	Welds and Turmeric	Reseda luteola L., Curcuma longa L.	Whole plant, Rhizomes	Yellow
83	Sapan wood and Madder	Caesalpinia sappan L., Rubia cordifolia L.	Wood, Root	Red
84	Madder, Turmeric and Welds	Rubia cordifolia L., Curcuma longa L., Reseda luteola L.	Root, Rhizomes, Whole plant	Orange, Red, Yellow
85	Madder, Berberry Wood and Logwood	Rubia cordifolia L., Berberis lyceum Royle/Berberis tinctoria Lesch., Haematoxylum campechianum L.	Root, extract	Brown
86	Ukulbir	Datisca cannabina L.	Wood	Brown, Yellow
87	Fir	Abies Mill.	Bark	Brown
88	Rasun Buti	Probably Allium sativum	Not known	Brown, Off-white
89	s.n.	Leucus sp.	Bark	Brown
90	Kauna	s.n.	Bark	Brown, Yellow, Off- white
91	Amlia sak	Amaranthus sp./ Hibiscus sp.	Not known	Ochre, White, Brown, Yellow, Yellow-ochre
92	Hinro	Scientific name undetected	Not known	Brown, Moss green, Red
93	Sakra beu	Scientific name undetected	Bark	Brown, Yellow, Off- white
94	Esfurkee patta	Delphinium sp.	Not known	Yellow
95	Ragira	Amaranthus sp.	Plants and Leaves	Yellow
96	Tamarix galls	Tamarix sp.	Galls	Black





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97	Cochineal	Dactylopius coccus Costa	Not known	Red, Yellow, Pink, Orange, Violet, Peach, Maroon, Bluish grey, Brown, Blue, Purple, Magenta
98	Lac	Scarlet resinous secretion of Laccifer lacca	Not known	Orange, Red, Maroon, Violet, Purple, Peach, Yellow-red, Brown, Pink
99	Haematin	Dark brownish pigment containing iron in the ferric state	Not known	Blue, Grey, Brown, Violet, Green, Purple
100	Lac and Indigo	Laccifer lacca, Indigofera tinctoria L.	Whole plant	Red, Orange
101	Indigo and Cochineal	Indigofera tinctoria L., Dactylopius coccus Costa	Whole plant	Grey, Blue, Red, Purple, Brown, Maroon, Pink
102	Madder and Cochineal	Rubia cordifolia L., Dactylopius coccus Costa	Root	Brown, Purple, Maroon, Red, Blue, Violet, Orange
103	Cochineal and extract of Jackfruits wood	Dactylopius coccus Costa, Artocarpus integrifolia L.	Not known	Yellow, Pink
104	Munjeet and Cochineal	Rubia munjith Roxb. ex Fleming, Dactylopius coccus Costa	Root	Red
105	Lac and Turmeric	Laccifer lacca, Curcuma longa L.	Rhizomes	Orange
106	Haematin, Barwood	Baphia nitida Lodd.	Not known	Grey
107	Cochineal, Pista flower and Sapan Wood	Dactylopius coccus Costa, Pistacia vera, Caesalpinia sappan L.	Flowers	Red
108	Cochineal, Madder and Turmeric	Dactylopius coccus Costa, Rubia cordifolia L., Curcuma longa L.	Root, Rhizomes	Orange
109	Lac, Cochineal and Indigo	Laccifer lacca, Dactylopius coccus, Indigofera tinctoria L.	Whole plant	Red
110	Cochineal, Berberry Wood and Logwood	Dactylopius coccus Costa, Berberis lyceum Royle/Berberis tinctoria Lesch., Haematoxylum campechianum L.	Root	Red

CONCLUSION

The present paper was an attempt to bring out or highlight the hundred and thirty-six year old legacy of Indian natural dyes which were very popular, rich and for which the India is famous. These hundred years old information can be helpful or a source of guide to all the people who is in interested in Indian dyes or plant based dyes. Many textile designers can be benefitted with these information by utilizing it for their customers who are drawn towards organic plant based dyes.

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RESEARCH ARTICLE

Optimum Solution on Two Stage Flow Shop Scheduling Model with **Identical Parallel Machines**

Khushboo Malhotra^{1,2}, Deepak Gupta², Sonia Goel^{2*} and A. K. Tripathi²

¹Department of Mathematics, D. A. V. College (Lahore), Ambala City.

²Department of Mathematics, Maharishi Markandeshwar Engineering College, MM (Deemed to be University), Mullana, Ambala.

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*Address for Correspondence Sonia Goel

Department of Mathematics, Maharishi Markandeshwar Engineering College, MM (Deemed to be University), Mullana, Ambala. Email: sonia.mangla14@gmail.com



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ABSTRACT

Efficient scheduling in manufacturing is crucial for optimizing production processes and minimizing costs. The type of FSSP considered in this study presents a challenging scenario. The present study gives a novel BB algorithm tailored for large problem instances. The proposed BB algorithm leverages the power of mathematical programming and MATLAB implementation to address large-scale instances of the problem, a critical need in contemporary manufacturing environments. This research aims to guide practitioners and researchers in selecting the most suitable method for optimizing make-span in manufacturing environments. The findings contribute to the ongoing efforts in the field of operations research and production scheduling. This study develops a Model for optimizing the elapsed time with two machines involving m identical parallel processors at first level and single processor at second level having utilization time of each task on each of the two machines. A MATLAB Code in R2018a is developed for introducing the proposed algorithm for very large number of tasks and machines. Numerical experiment has also been done for supporting the coding.

Keywords: Scheduling, Mathematical Model, MATLAB, Identical Parallel Machines.





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INTRODUCTION

Efficient production scheduling is a cornerstone of modern manufacturing, as it directly impacts productivity, cost-effectiveness, and overall operational performance. Among the various scheduling problems that industries face, this problem which is considered in this study presents a complex and practically significant challenge. This scheduling scenario is particularly common in industries where optimizing the sequential processing of jobs through different stages is paramount. In this paper, we embark on a comprehensive exploration of scheduling techniques to address the FSSP with a focus on the specific configuration of identical parallel machines. The primary objective is to minimize the make-span, the total time required to complete all jobs, while respecting the constraints inherent to this problem. This study evaluates a novel Branch and Bound algorithm, specifically designed to tackle large problem instances. Leveraging the power of mathematical programming and implemented in MATLAB, this algorithm aims to provide an efficient solution to the scheduling problem, especially pertinent for manufacturing environments where the scale of the problem is substantial.

PRACTICAL SITUATION

The real life application of our model can be shown through the following illustration:

Pharmaceutical Manufacturing

Stage 1 (Parallel Machines) Multiple machines can be used for various stages of pharmaceutical tablet manufacturing, including mixing, granulation, and compression.

Stage 2 (Single Machine) A single machine, like a tablet coating machine, can be used in the second stage for coating tablets with the desired coating material.

Here the two-stage flow shop model is applied to optimize production processes, enhance quality control, and manage the flow of materials and products efficiently. The use of parallel machines in the first stage improves throughput, while the single machine in the second stage ensures precision and consistency in critical operations.

LITERATURE REVIEW

Efficient scheduling in manufacturing operations is vital to enhance productivity and reduce costs. Next, if Scheduling is done with the help of parallel machines then it would be a boon for manufacturing field. Hence in this study we are considering parallel processors. Parallel machine scheduling (PMS) means to perform tasks on multiple processors. Feifeng Zheng also did work in this field (Zheng, 2022). Malhotra et. al. also worked on FSSM with identical parallel processors (Malhotra, 2022). Many methodologies have been considered for obtaining performance measures.But here we are considering branch and bound in this paper. Branch and Bound algorithm is a classical technique often applied to various scheduling problems. In the context of flow shop scheduling, it has shown its potential for solving problems with smaller instances (Sullivan &Daganzo, 1986). Many researchers have worked on this BB method like (Gupta D., 2022), (Gupta D., 2020), (Malhotra K., 2022), (Malhotra K., 2023) and many more. Researchers have explored variations of the two-stage flow shop scheduling problem, including considering job release times, due dates, and machine breakdowns. Hazzi et. al. also worked on two stage flow shop scheduling model (Hazzi, 2023). Liu and Pardalos also worked with parallel machines (Liu, 2023). Hnaien also studied on two stage flow shop scheduling model (Hnaien, 2023). A comprehensive survey by Hino and Morabito (2017) provides insights into different scheduling problems and their solution methods. Jemmali et. al. worked on two stage flow shop scheduling model (Jemmali , 2023). Parviznejad also studied the problem related to parallel machines (Parviznejad, 2021). Hidri also worked on parallel machine scheduling (Hidri, 2022).

PROBLEM DISCRIPTION

Let us assume that there are n jobs for performing on two machines E and F. Type E has m identical processors and Type F has single processor where ei and fi denote consumption time of task i on E and F processors respectively. It is not necessary to perform each job on all parallel machines of type E, one job can be performed partially on parallel





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machines of type E and then after processing on first processor, it will be performed on second processor. Offered time of all similar processors t1j are also given. The purpose of the problem is to manage scheduling in such a way to complete project in minimum time. The model is represented as:

MATRIX FORM OF MODEL

The model of the problem is depicted in matrix form is given in Table 1 which is given below:

SOLUTION APPROACH PROPOSED METHODOLOGY

We solve this problem as follows:

Step 1:

Now check the condition

$$\sum_{j=1}^{m} t_{1j} = \sum_{j=1}^{n} e_{i}$$
 (1)

and apply MODI method otherwise make arrangement to satisfy condition (1).

Step 2:

Apply the formula

$$1' = \max_{i \in Jr} \left(\sum_{i=1}^{n} e_{ij} \right) + \min_{i \in Jr'} (fi)$$
(2)

$$1" = \max_{i \in Jr} \left(e_{ij} \right) + \sum_{i=1}^{n} (fi)$$
 (3)

- a. Find $1 = \max\{1', 1''\}$
- b. Extract out I for all arrangement of tasks.
- c. After that take minimum value from all l's. This will prepare branching tree.
- d. In last, we have to generate In-Out table for the required sequence of tasks.

NUMERICAL APPROACH

The numerical problem is given in Table 2.

SOLUTION

We solved the problem in four Steps.

Step 1: See MODI method in Table 3:

Step 2: Apply branch and bound method and get the final table

The final In-Out Table of required sequence is given in Table 4. The above gantt chart is showing final In-Out table with six parallel machines of type E and single machine of type F. Here minimum make-span is 37.6620 hrs. Hence, the problems of small size where either the no. of jobs are less or the no. of equipotential parallel machines are less are easily solvable manually but for large sized problems we have generated MATLAB program in R2018a version for finding make-span for large sized problems. Table 5 is shown by taking more parallel machines at level 1 and more no of jobs also.

CONCLUSION

This study addresses the critical issue of efficient scheduling in manufacturing, focusing on the Two-Stage Flow Shop Scheduling problem with m identical parallel processors at the first level and a single processor at the second level. The development of a novel Branch and Bound algorithm specifically tailored for large-scale instances of the problem sets this research apart. The algorithm's implementation in MATLAB 2018a enables its application to real-





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world manufacturing scenarios, where optimizing the make-span is essential for minimizing costs and improving production processes. The proposed Branch and Bound algorithm demonstrates its efficacy in handling large problem instances, showcasing the power of mathematical programming in addressing complex scheduling challenges. The numerical experiments conducted not only validate the efficiency of the proposed algorithm. The findings offer valuable insights into the strengths and weaknesses of each scheduling approach, guiding both practitioners and researchers in selecting the most suitable method for optimizing Two-Stage Flow Shop Scheduling in manufacturing environments. This research contributes to the broader field of operations research and production scheduling, enhancing our understanding of how diverse techniques can be employed to bolster manufacturing competitiveness. Overall, the study underscores the significance of tailored algorithmic approaches in addressing specific scheduling challenges within the manufacturing domain.

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Table 1. Matrix Form

Jobs	Machine E		Machine F	
I	E ₁ E ₂ E ₃ E ₄ E ₅	Em	Utilization Time (ei)	Utilization Time (fi)
(1)	e_{11} e_{12} e_{13} e_{14} e_{15}	$e_{1m} \\$	e ₁	f ₁
(2)	e21 e22 e23 e24 e25	e _{2m}	e 2	f ₂
(3)	$e_{31} e_{32} e_{33} e_{34} e_{35} \dots$	e_{3m}	es	f ₃
•			•	
			•	
			•	
(n)	$e_{n1}e_{n2}e_{n3}$ e_{n4} e_{n5}	e_{nm}	e n	fn
Available	$t_{11}t_{12}$ t_{13} t_{14} t_{15} t_{15}	•		
Time	111 112 113 114 115 1	lm		

Table 2. Defined problem

Jobs	E 1	E 2	Е3	E4	E 5	E 6	Processing time (ei)	Processing time(fi)
1	1	6	6	6	7	7	3.666	3.333
2	5	6	7	8	9	9	2.666	6
3	2	2	2	3	3	2	3	5.333
4	5	6	7	8	9	9	4.333	5.333
5	2	2	3	3	3	2	4.666	6
6	5	6	7	8	9	9	3.666	4.666
7	5	6	7	8	9	9	4.333	5.333
t _{1j}	9	9	8	7	6	5		

Table 3. Modi Method

Jobs	E ₁	E 2	Ез	E ₄	E 5	E 6	F
1	3.6660	0	0	0	0	0	3.333
2	2.6660	0	0	0	0	0	6.000
3	0	0	3.0000	0	0	0	5.333
4	0	4.3330	0	0	0	0	5.333
5	0	0	0	0	0	4.6660	6.000
6	0	1.6640	1.0010	0	0	0	4.666
7	0	0	4.3330	0	0	0	5.333
8	2.6680	1.0010	0.6670	7.0000	6.0000	0.3340	0

Table 4. In – Out table for best possible sequence

Jobs	E 1	E ₂	Ез	E4	E 5	E 6	F
6	-	0 - 1.6640	0-1.0010	-	-	-	1.6640-6.3300
2	0-2.666	-	-	-	-	-	6.3300-12.3300
8	2.666-5.334	1.664-2.665	1.0010-1.6680	0-7	0-6	0-0.3340	12.3300-12.3300
7	-	1	1.6680-6.0010	-	ı	1	12.3300-17.6630
5	-	1	-	-	ı	0.3340-5.000	17.630-23.6630
4	-	2.6650-6.9980	-	-	ı	1	23.630-28.6630
3	-	-	5.0010-9.0010	-	-	-	28.6630-34.3290
1	5.3340-9.0000	-	-	-	-	-	34.3290-37.6620

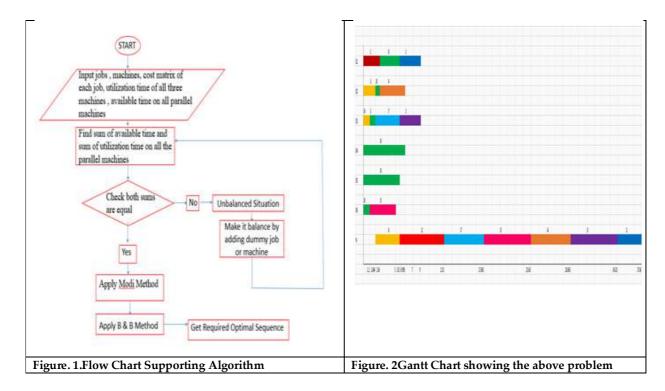




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Table 5. Matlab output table for proposed Algorithm

Number of Tasks	Number of Processors at level 1	Optimal Sequence of Jobs	Optimum Make span in Hrs.
4	6	4,3,5,2,1	26.6650
5	6	2,5,6,4,3,1	36.6650
7	6	6,2,8,7,5,4,3,1	37.6620
10	6	1,2,11,10,9,8,7,6,5,3,4	51.6630
15	6	9,10,2,13,12,15,1,14,7,3,5,6,8,11,4	115.9997
20	6	10,1,11,2,20,15,17,9,14,16,4,5,8,12,19,7,10,6,13,3	137.6583
30	6	26,24,1,11,18,17,29,28,20,12,15,9,16,2,4,5,8,19,23 27,30,7,10,22,25,14,6,21,13,3	195.6600
50	6	46,49,1,45,42,48,50,9,47,44,43,11,16,24,26,31,39,41,2,4,5,8, 12,15,17,19,20,23,27,30,32,34,35,38,7,10,22,25,37,40,14,29,6,21 36,3,13,18,33,28	328.6550
80	6	9,14,1,12,11,13,80,10,15,2,79,16,24,26,31,39,41,46,54,56,62,67,75 77,4,5,8,17,19,20,23,27,30,32,34,35,38,42,45,47,49,50,53,57,60,63 66,68,70,71,74,78,7,22,25,37,40,52,55,61,73,76,29,44,59,65,6,21 36,51,72,3,18,28,33,43,48,58,69,64	522.6420







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Trends and Future Applications for Environmental Remediation

RESEARCH ARTICLE

Insights into the Photo Catalytic Degradation of Phthalates: Current

Urvashi, Gunjan Chauhan*

Department of Chemistry, MMEC, Maharishi Markandeshwar (Deemed to be University), Mullana, Ambala Haryana 133207.

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*Address for Correspondence

Gunjan Chauhan

Department of Chemistry,

MMEC, Maharishi Markandeshwar (Deemed to be University),

Mullana, Ambala Haryana 133207.

Email: gunjan.chauhan@mmumullana.org

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ABSTRACT

Esters of 1, 2-benzenedicarboxylic acid commonly known as phthalates have a wide variety of industrial, agricultural and many other applications. The most significant usage of phthalates is as plasticizers, i.e. increase the functionality and flexibility of polymeric materials. However, researches have shown that exposure to phthalates can lead to detrimental health effects, such as hormone disruption, increased risk of asthma and allergy, type II diabetes, insulin resistance, reproductive and developmental issues. This raises concerns about the safety and sustainability of their widespread use in almost every industry. Numerous regulatory bodies classified phthalates, a top priority pollutant, mandating the reduction and control of "phthalate pollution". The US Environmental Protection Agency (EPA) and European Union(EU) implemented regulations that require companies to monitor and limit the use of phthalates in consumer products, also prompting manufacturers to reformulate their products to reduce phthalate content and ensure compliance. The present study provides a clear perspective on the recent progress in the degradation of phthalates by photo catalytic degradation, also highlighting its potential for environmental remediation. Various researches have focused on exploring semiconductor-based photo catalysts, and their modifications to enhance degradation efficiency. Future prospects encompass the fabrication of novel photo catalysts, mechanistic understanding, scaling up processes for practical applications, also addressing challenges towards sustainable and efficient phthalates degradation.

Keywords: phthalate pollution, hormone disruption, development issues, photo catalytic degradation, semiconductor-based photo catalysts (SBP)





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INTRODUCTION

The manufacturing, consumption, and disposal of plastics have resulted in the release of phthalates in environmental matrices, including the air, groundwater, sediments, waste water, streams and lakes [1]. Phthalates, which are also known as phthalic acid esters such as di(2-ethylhexyl) phthalate (DEHP), diisononyl phthalate (DINP), diisodecyl phthalate (DIDP), and di(2-propylheptyl) phthalate (DPHP) are used in poly vinyl chloride compounds (PVCs) to increase the flexibility and stability of plastics. In non-polyvinyl compounds (non-PVCs), phthalates such as dimethyl phthalate (DMP), diethyl phthalate (DEP), butyl benzyl phthalate (BBP), and di-butyl phthalate (DBP) used to improve the stability and solubility in cosmetic products like hair growth sprays, perfumes, nail polishes, cleansers, and shampoos[2-4]. The plasticizers are frequently utilized as phthalates, readily and extensively released into the environment through a variety of processes, such as leaching, migration, evaporation during product use, the disposal of industrial and municipal solid wastes[5]. Environmental concerns have been raised due to the hepatotoxic, teratogenic, and carcinogenic properties of phthalates, which are caused by the large manufacturing units and usage of phthalate containing products[6].

The phthalates are used often to soften plastics in a variety of consumer goods, including office supplies, footwear, flooring, toys, coated paper and fabrics, recreational equipment, beds, cables and others[7]. The exposure of bottled water to direct sunlight and the interaction of hot water with disposable plastic and paper cups both significantly raised the levels of phthalate in the drinking water (p value < 0.05)[8-10]. A team of scientists and medical professionals with expertise in neurotoxicity and toxic chemicals, recently came to the conclusion that there is strong evidence connecting phthalate exposure to a higher risk of behavioural, attention, and learning issues in kids. It has been suggested that phthalates should be removed from goods that could expose vulnerable groups, such as pregnant women, and children[11]. There is now increased focus on developing technology that can efficiently achieve degradation and removal of these compounds. In this review, phthalate photo catalysis degradation is reviewed in depth to reduce environmental contamination. This review paper explores the use of semiconductor photo catalysis as a viable option for phthalate removal. An overview of photo catalyst development is provided which covers a broad range of materials that have been reported in the literature. In addition, an overview of the effects of phthalates on human health, including the mechanisms of phthalate degradation and risk assessment are reviewed to provide the reader with a detailed insight into the photo catalytic removal mechanism.

PHTHALATES SOURCES AND CATEGORY

Phthalates are used in the production of variety of consumer, industrial as well as personal care goods. Phthalates are generally utilized as plasticizers in the manufacturing industries to create (PVC). In the majority of cases, phthalates are added to the polymers to make them softer and more malleable[12]. From the past two decades the European Union (EU) and United state Environmental Protection Agency (USEPA) has prohibited the use of six phthalates due to their potential hazards to health. The phthalates include (DEHP), (BBP), (DBP), (DINP), (DIDP), and (DNOP)[13].

Categories and applications of phthalates

Tere- and ortho-phthalates are the two types of phthalates that tend to be used most frequently. Numerous types of alcohols, including methanol, ethanol, and tridecyl alcohol, react with phthalic anhydride to form ortho-phthalates, which are then simply referred to as "phthalates"[14]. The ortho-phthalates are classified into two categories: ortho-phthalates with a high molecular weight (HMW) such as DEHP, DNOP, DIBP, DIDP, and low molecular weight (LMW) include DMP, DEP, DBP listed in the table 1. The molecular structure of HMW phthalates has seven to thirteen carbon atoms, while LMW phthalates has three to six carbon atoms only [15].





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Exposure routes of phthalates Through food

Food is presumed to be the primary source of exposure and the majority of the population is exposed to phthalates through dietary supplements thus they have received increased attention. Phthalate regulations have been established for plastic packaging materials related to food, with limits of 1.5, 0.3, and 30 mg/kg for DEHP, DBP, and BBP, respectively[16]. Edwards et al. carried out a preliminary investigation in order to examine the amounts of ortho-phthalate and replacement plasticizer in food and food handling gloves from fast food restaurants in the United States. The highest quantities of DEHT were detected in gloves. The results indicated that the food samples consist of 70% DEHP and 81% DnBP. The average DEHT concentrations in burritos were much higher than in hamburgers (6000 μ g/kg vs. 2200 μ g/kg; p < 0.0001)[17].

Through air

Phthalate concentrations in indoor air are frequently high but can also vary based on the presence of phthalate[18]. Huang et al. conducted a study by taking gas and PM2.5concentrations of 7 phthalates from 40 residences were measured during summer and winter. Higher air humidity led to more water absorption of aerosols in summer, facilitated mass transfer of phthalates from air to PM2.5, and resulted in greater Kp of DiBP and DnBP in the summer. According to the results, the overall amounts of airborne phthalates (gas PM2.5) significantly dropped from summer to winter[19].

Through water

Hot water in disposable plastic and paper cups raise the levels of phthalates in drinking water.DMP and DEHP were the main phthalate-containing chemicals, contributing more than 60% of the overall phthalate levels across all water sources[20]. Tehran, Spain, China and Iran, Abtahi et al. investigated the health risks associated with phthalate exposure from drinking water, and they also examined the presence of phthalates in bottled, tap, and water resources. The results showed that the average concentrations of total phthalates (\pm standard deviation: SD) were 0.76 \pm 0.19, 0.96 \pm 0.10, 1.06 \pm 0.23, and 0.77 \pm 0.06 μ g/L in surface waters, ground waters, bottled water, and drinking water from the water distribution system, respectively. DMP and DEHP were the leading phthalate-containing chemicals, contributing more than 60% of the overall phthalate levels in all water sources[4].

Toxicity effects

The widespread use of plastics associated with the use of variety of substances such as additives and phthalates chemicals that are discharged into the environment, are among the primary effects of industrialization[21]. In 2021, an astounding 390.7 million metric tones of plastic has been manufactured globally, of which 9% was recycled, 12% was burned, and the remaining was disposed of directly into the landfills or the environment[22]. Numerous of these substances bio accumulate in biological tissues and can have a deleterious effect on the health of people and animals by changing endocrine functions. Therefore, through skin contact, ingestion, or inhalation of these widely dispersed environmental toxicants, living organisms including people and animals are invariably exposed to endocrine disrupting chemicals (EDCs)[21]. The hormone Society and the International Pollutants Elimination Network (IPEN) addressed the growing concern in the economy regarding the numerous health effects of hormone disruptive chemicals found in phthalates[23].

Toxic Effect on Aquatic Organisms

The harmful effects of phthalates concentrations are dependent on the type and toxicity of the phthalate. Phthalates are hazardous to the immune system, endocrine system, metabolism, and produce oxidative stress in the aquatic organisms. Exposure to phthalates in adult aquatic organisms causes damage to the liver, kidney, reproductive system, and other organs. It has been determined via numerous studies that phthalates exposure to aquatic organisms pose both acute and long-term health effects. Cong *et al.* conducted a study to examine the effects of (DMP) on adult zebrafish (*Danio rerio*). The research findings shows that the survival is dependent on the DMP dosage. Within 96 hours, only 20% of the fish died in the presence of 25 mg/L of DMP, while the remaining fish





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managed to survive. 80% of the fish died with 100 mg/L DMP died within 96hours. In the 200 mg/L concentration of DMP, fatality happened in 96 hours at a rate of 100% [24] . Furthermore, exposure to DEP, DBP, BBP, and DEHP produce neurotoxicity by affecting the activity of genes present in aquatic organisms such as mRNA (messenger ribonucleic acid) and acetylcholinesterase (AchE). Chen et al. studied the effects of exposure to the different concentrations of Di-n-butyl phthalate (DBP) (20 μ g L⁻¹, 100 μ g L⁻¹, and 500 μ g L⁻¹) on juvenile red snapper Lutjanus argentimaculatus, for a time period of 15 days. The study demonstrated that fish exposed to DBP shows that the ethoxyresorufin O-deethylase (EROD) activities and cytochrome P4501A (CYP1A) levels in the liver and lungs is dependent on time, concentration, and tissue [25].

Toxic Effects on Human Health

Exposure to phthalates with human body through the mouth, nose, or skin cause detrimental health effects. Both the male and female reproductive system is adversely affected. A Chinese study examine that the exposure to the phthalates metabolites: monomethyl phthalate (MMP), monoethyl phthalate (MEP), monobutyl phthalate (MBP), monobenzyl phthalate (MBZP), mono-2-ethylhexyl phthalate (MEHP), and mono-2-ethyl-5-oxohexyl phthalate (MEOHP) strongly impact the male semen quality [26, 27]. Among the several types of phthalates, DEHP is the most commonly used and hazardous. Also has the potential to negatively affect human health and functions as an endocrine disruptor [28]. Another prospective study demonstrated increased phthalate concentrations will decrease the oocyte fertilization, mature oocyte and good-quality embryos. Thus, it appears that exposure to phthalates is associated with early puberty, preterm birth, and certain conditions that impact female fertility, including endometriosis and DMP. Furthermore, the unfavourable outcomes male foetuses face during pregnancy may be due to phthalate's exposure.

PHOTOCATALYSIS MECHANISM FOR SUSTAINABLE DEGRADATION OF PHTHALATES

The term "photo catalyst" refers to a semiconductor material that can accelerate a chemical reaction by the absorption of photons of energy (in ultraviolet or visible region) greater than or equal to 3.2 eV. The photochemical process initiates when photons with energy greater than or equal to the photo catalyst material ($hv \ge E_{gap}$) interacts with the surface of photo catalyst[29]. The simultaneous photochemical redox reactions can take place. As a result of this photo redox reaction the electron-hole (e⁻/h⁺) pairs are generated in the conduction(e^-_{cb}) and valence bands(h^+_{vb}) (equation) respectively[30]. The produced e^-_{cb} and h^+_{vb} are strong reducing and oxidizing agents[31]. The e^-_{cb} combine with oxygen molecule to form superoxide radical (O^-_2) whereas the h^+_{vb} combine with water molecule to produce hydroxyl (OH-). The produced (OH-) with strong oxidation potential of 2.06 eV degrades pollutant molecule(X) such as phthalates, irrespective of the X selectivity. The pollutant molecule degradation mechanism is mentioned in equation (a-d)

$$Photocatalyst + hv \ge Egap \rightarrow h_{vb}^{+} + e_{cb}^{-}$$
 (a)

$$h_{vb}^+ + H_2O \rightarrow Photocatalyst + OH + H^+$$
 (b)

$$e_{cb}^{-} + O_2 \rightarrow Photocatalyst + O_2^{-}$$
 (c)

$$OH + X \rightarrow intermediates \rightarrow CO_2 + H_2O + by products$$
 (d)

Semiconductor-based photo catalysis

The semiconductor photo catalysts have drawn more attention in recent years, due to potential applications in the environmental purification and mitigation of organic as well as inorganic pollutants from water and air [32]. The photo catalytical degradation is considered to be one of the best photochemical processes for mineralisation of the personal care products (PCPs) particularly phthalates. In 2022, Pang et al. fabricated two commercial catalysts TiO₂, ZnO for the degradation of the Diisobutyl phthalate (DiBP). The results show that greater than 90% of degradation takes place within 25 min. The generated super oxide radical •OH and •O2 are essential components in the photo catalytic degradation of DiBP for both materials. This is the first research on semiconductor photo catalysis-induced phthalate DiBP disintegrate in water[33]. The alkaline earth metals are abundant and inexpensive in the earth's crust, their stannate perovskite oxides offer a wider range of potential applications in multiple fields. The band gap value, friendliness affordability, stability, and environmental MgSnO3 and CaSnO3





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which enable them to degrade phthalates. Ye et al. investigated the photocatalytic degradation performances of dimethyl phthalate (DMP) and diethyl phthalate (DEP) by the oxides of stannate perovskite MgSnO3 and CaSnO3under simulated sunlight. The photochemical examination results demonstrated that MgSnO3 perovskite performed better when compared to CaSnO₃ perovskite. This is due to increased specific surface area, pore volume, average pore width, greater abundance of oxygen vacancies which facilitated the adsorption, transport, and activation of reactive oxygen species [34]. According to reports, PAEs that have an alkyl chain length of less than six such as DMP, DEP, and DBP are intrinsically harmful to aquatic life. Thus allows them to be easily absorbed by the skin and digestive system, and then subsequently transmitted to the majority of the body tissues, including the placenta[35]. Wang et al.'s research illustrated that phthalic acid esters dimethyl phthalate (DMP), diethyl phthalate (DEP), dibutyl phthalate (DBP) undergoes photo catalysis degradation by the UV, UV/TiO2, and UV-Vis/Bi2WO6. The results indicate that, the UV/TiO2 exhibited the highest degrading efficacy on a particular group of PAEs, achieving removal efficiencies of up to 93.03, 92.64, and 92.50% in 90 minutes for DMP, DEP, and DBP, respectively. However, the system UV-Vis/Bi₂WO₆ was completely inefficient for eliminating DEP and DMP[36]. Zhang et al. synthesized a Z-scheme nanocomposite rGO-(Bi₂O₃-TiO₂) for the removal of the Di(2-ethylhexyl) phthalate (DEHP) is a known carcinogen, teratogen, and mutagenic environmental contaminant. The findings show that, doping a composite Bi₂O₃-TiO₂ with 4% rGO, shows the best photo catalytic degradation of 89% within 90 minutes of continuous solar light irradiation. By considering the reaction parameters such as pH=6 of the solution and concentration of DEHP (10 mg L-1). Therefore, the created hetero junction enhanced electron-hole separation rate in [rGO-(Bi₂O₃-TiO₂)]. Furthermore, the photo catalytic performance of Bi₂O₃-TiO₂ was 1.61 times stronger than that of pristine Bi₂O₃ and TiO₂.[37].

Abbreviations

DMP (dimethyl phthalate), DEP(Diethyl phthalate), DBP (Dibutyl Phthalate), DEHP (Di-Ethyl-Hexyl-Phthalate), BBP(Butyl-Benzyl-Phthalate), DIBP (Diisobutyl phthalate)

CONCLUSION

Phthalates are one of the categories of emergent contaminants that have the greatest negative environmental impact due to their extensive use as well as constantly spilled into the reservoirs. The origins of phthalate contamination, its health effects, the exposure routes, and, photo catalytic mineralisation technique have all been covered in this analysis. This study provided a comprehensive summary of the disrupting-effects of phthalates on living organisms such as the human and animals. Currently, the effects of phthalates on aquatic organisms are better documented than the effects on human health. Therefore, there is need to evaluate this issue better and improve the negative impact of exposure to phthalates on the both vivo and vitro. The degradation of the phthalate's contaminants by advanced oxidation processes such as photo catalysis is important for environmental remediation. The review aims to provide information on how photo catalytic process has become a rapid and important strategy to mitigate the pollutants released from the overuse of the phthalates containing products.

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Table 1 displays the categories, molecular formula, photocatalyst, % degradation and time period, experimental conditions, and uses of the phthalates

Compounds	Molecular Formula	Photocatalyst	% degradation and time period(min)	Experimental conditions	Uses	Ref
Dimethyl phthalate (DMP)	C10H10O4	N/TiO2 and N/TiO2	N/TiO ₂ (41%) and UN/TiO ₂ (58%) within 5 hrs	DMP initial concentration: 15 mg/L Temperature: 22±2 °C Catalyst loading: 0.5 g/L.	Used in hair sprays, enteric coatings for medicines and rubber, solid rocket propellants, nail polish, artificial nails, insect repellents, and pesticides.	[38]
	C10H10O4	20 wt % Bi ₂ WO ₆ /ZnO (20-BWZ)	86.6% within 90 min	DMP degradation under natural sunlight irradiation		[39]
Diethyl phthalate (DEP)	C12H14O4	0.2 mol % of Zn doped TiO ₂	100% within 200 min	mercury lamp =125 W catalyst load = 1 g/l. Zn doped TiO ₂ calcined at 450 °C.	Directly added in cosmetic products and indirectly in fragrances. Also used as a solvent and alcohol denaturant.	[40]
		TiO ₂ (anatase)	100% within 50 mins	initial concentration: 20 mg/L Catalyst loading: 0.2 g/L Temperature=25°C pH=4.0 Luminous intensity: 2.5×10-6 Einstein/L·s.		[41]
		ZnO/SiC	90%within 120 and under visible light irradiation the time for maximum degradation was nearly 660 min.	DEP initial concentration: 5, 10, 15 and 25 mg/L pH: ranging from 2 to 12.		[42]
Dibutyl Phthalate (DBP)	C16H22O4	3% Ce/Gd- WS2	75% within 60 min	initial concentration of catalyst = 5ppm pH= 7 catalyst loading =10 mg	Plastics such as PVC, adhesives, printing inks, sealants, grouting agents, perfumes, deodorants, hair sprays, nail polish, and insecticides.	[43]
		(Fe, Ag) co-	95% within	DBP initial		[44]





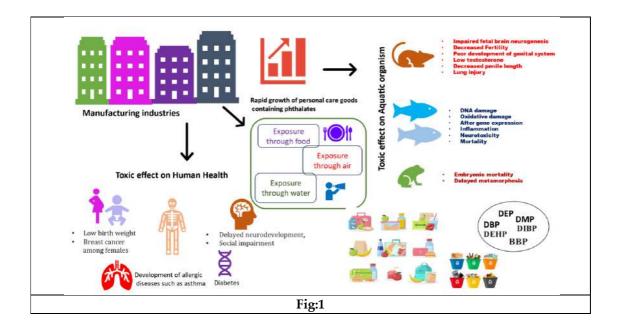
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		doped ZnO	137.5 min	concentration: 15 mg/L pH: 3 Catalyst loading: 0.15 g/L.		
Di-Ethyl- Hexyl- Phthalate (DEHP)	C24H38O4	rGO-(Bi ₂ O ₃ – TiO ₂)	89% within 95 min	pH = 6 Concentration of DEHP =10 mg L ⁻¹ reaction temperature =25 °C catalyst dosage = 0.0834 mg L ⁻¹	Diapers, food containers, plastic film for food packaging, blood bags, catheters, gloves, and medical equipment's such as tubes for fluids etc.	[37]
		Fe-Ag@ ZnO	90 % within 120 min	Catalyst loading: 20 mg/L. The authors reported that photocatalytic degradation of DEHP increased when operated at a higher and lower pH (as a result of acidic and basic catalytic hydrolysis).		[45]
Butyl- Benzyl- Phthalate (BBP)	C19H20O4	TiO ₂ /UV	70% and 120 min	catalyst dosage = 2.0 g L ⁻¹ pH=7.0 concentration of BBP = 1 mg L-1 concentrations of the co-existing substances = 0.01 M	Perfumes, hair sprays, adhesives and glues, automotive products, vinyl floor coverings	[46]
Diisobutyl phthalate (DIBP)	C16H22O4	P25 TiO ₂ and ZnO	>90% and 30 min	catalyst loading = 20 mg concentration of DiBP = 10 mg L ⁻¹	utilised as plasticizer in solid rocket propellants, nitrocellulose lacquers, elastomers, explosives, printing inks, adhesives and nail polish.	[33]





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RESEARCH ARTICLE

Preparation of Graphene Oxide: Various methods and their comparison

Ajay Jakhar and Jyoti Sharma*

Department of Chemistry, MMEC, Maharishi Markandeshwar (Deemed to be University), Mullana-133207 (Ambala) Haryana, India.

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*Address for Correspondence Jyoti Sharma

Associate professor, Department of Chemistry, MMEC, Maharishi Markandeshwar (Deemed to be University), Mullana-133207 (Ambala) Haryana, India. Email: jsharma117@gmail.com



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ABSTRACT

Graphene oxide (GO), a versatile and extensively studied material, finds applications in a myriad of fields owing to its unique properties. This abstract provides a comprehensive overview of various methods for the preparation of graphene oxide, drawing insights from seminal and contemporary research. The Hummers' method, initially introduced in 1958, involves the oxidation of graphite using a mixture of potassium permanganate, sulfuric acid, and sodium nitrate. Brodie's method, a historical precursor to Hummers', employs nitric acid as the oxidizing agent. Building upon these foundational approaches, the Improved Hummers' method and chemical exfoliation of graphite have been proposed, allowing for adjustments in reaction conditions and alternative oxidants. Ultrasound-assisted methods utilize acoustic energy for exfoliation, while microwave-assisted exfoliation accelerates the process through targeted irradiation. Electrochemical exfoliation, on the other hand, involves the reduction of graphene oxide on an electrode. The review synthesizes these methods, highlighting their respective advantages and limitations, providing a valuable resource for researchers engaged in the synthesis of graphene oxide for diverse applications. As the field continues to evolve, this overview serves as a foundation for understanding and exploring novel techniques for graphene oxide preparation.

Keywords: Graphene Oxide, Hummers method, Exfoliation methods, Oxidizing agent and Microwaveassisted methods.





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INTRODUCTION

When in 1859 Graphene oxide was synthesized by Brodie[1], there is a surge in its research after the introduction of electronic properties of Graphene by Giem and Novoselov[2] as it has a prominent role in the development of Graphene. On investigating the nomenclatures of Graphene and graphene oxide they show some similarities as they both are carbon nanomaterials howsoever graphene and GO are very different, Graphene is a single layer of SP2 hybridized Carbon atoms whereas GO has a carbon structure with defects and oxygen containing functional groups which imparts to its SP3 hybridization to a substantial degree. The presence of oxygenated groups in GO affects its electronic, mechanical and electrochemical properties. These polar functional groups of GO makes it strongly hydrophilic[3] making it easy for GO to link to other metals and nonmetallic nanoparticles by noncovalent or covalent chemical bonds which creates extraordinary optical and electrical properties for GO based nanocomposites[4,5] and stable GO dispersion for preparation of thin conductive films by drop-casting or spin coating[6]. The structure of GO and positioning of GO is been a topic of debate ever since and exact structure is yet to be found though the Lerf-Klinowski[7] model is been widely accepted Fig-1. In this model, Hydroxyl and epoxide functional groups are proposed to decorate the basal plane which are segregated into islands among the lightly oxidized, graphene-like regions while carboxylic acids or carbozlates, depending on the pH of the solution are present on the edges of the sheets. Electronic properties of GO sheets depend on degree of structural disorder from the presence of substantial amount of sp³carbon fraction. GO films have a energy fap in the electron density making it typically insulating in its natural state[8], in sheet form GO exhibits high sheet resistance (R_s) about 10¹² \Omega sq⁻¹[9]. GO also exhibits excellent photoluminescence[10] which occur from near-UV-to blue visible (vis) to near-infrared (IR) wavelength range which makes GO very useful in biosensing, optoelectronics applications.[11]

Electrochemical properties of GO are also fascinating as because of its favorable electron mobility and unique surface properties, such as one-atom thickness and high surface area, GO can accommodate the active species and facilitate their electron transfer at electrode surfaces as reported by Zuo et al[12], GO supports the efficient electrical wiring of the redox xenters of several heme-containing metalloproteins to the electrodes. Graphene oxide is an excellent candidate for fluorescence resonance energy transfer sensors for single stranded DNA monitoring, GO has unique interaction with nucleic acid makes it simple to identify and quantify a target DNA.[13] Abundance of oxygen containing functional groups it binds to various catalysts for hydrogen storage and generation, taking advantage of bandgap of GO it has wide application iin LEDs having GO interlayers reaching a high value in luminosity of 220% increase and 280% power efficiency in comparison to devices without GO.[14] In a comparative study of oxidation synthesis methods of Graphene Oxide Pumera et al[15] concluded in their study that permanganate-based oxidations such as Hummers and Tour methods introduced a greater amount of oxygen functionality and higher proportion of carbonyl and carboxylic acid groups than methods that utilize chlorate-based oxidants such as Staudenmaier, the Brodie, Hofmann methods but chlorate based oxidations methods has more uniform functionality and less extensive methods and materials. In this review we try our best to summarise the abundantly used methods for preparation of Graphene Oxide- Hummers Method, Brodie Method, Staudenmaier Method, Electrochemical Exfoliation methods, Microwave Exfoliation methods.

Methods for GO preparation

Protocols and procedures used in various methods are summarized in a brief manner as follows

Hummers method

In Hummers' method concentrated H₂SO₄ behaves as the intercalator and KMnO₄ acts as the oxidant[16-17] and the ratio of these are very crucial for preparation of good quality of Graphene oxide. Procedure used by Hummers is briefed in Fig-2 Formation of Graphene oxide from graphite can be divided into three individual steps as explained by Dimiev[18], where each step can be stopped and each intermediate product can be isolated and stored at appropriate conditions. First step involves formation of graphite intercalation compound with H₂SO₄(H₂SO₄-GIC) which is a crucial step for GO formation as this step allows diffusion of oxidizing agents into the interlayer spaces of





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graphite, sodium nitrate has synergistic effect in the process. In second step GIC is converted to pristine oxidized graphite (PGO), this is rate determining step which takes hours to complete. In third step PGO is converted to GO with exposure with water. We can also modify the timing of reaction in Hummers' method for production of well exfoliated Graphene oxide as shown by Aixart[19], where they produced GO with reaction time as GO-30 min, GO-60 min, GO-120 min, GO-300 min, GO-540 min and concluded that with increase in reaction time and avoiding aggressive steps during end of reaction we can produce greater exfoliated GO with increased interlaminar spacing. Graphene oxide synthesized from Hummers' method contains higher amount of carbonyl and carboxyl groups when compared to GO from Brodies' method.

Brodies' Method

This method involves high vaccum conditions and samples are efficiently dried from residual water to yield correct C/O ratio (atomic percent).[20] Graphene oxide synthesized by Brodies' method[1](Procedure briefed in Fig-3) has high relative amount of epoxy and hydroxyl groups on the planar surface of graphene oxide flakes as compared to graphene oxide synthesized from Hummers' method. The GO produced from brodies method has as interlayer spacing of 7.0±0.35Å[21]and functional groups such as hydroxyl, epoxide, carboxyl and some alkyl groups are distributed randomly. Hydroxyl and epoxide groups are located on their basal planes close to each other and carboxyl and alkyl groups were located at the edges of the GO flakes, when characterised by EA, XRD, SEM, TEM, RAMAN, FTIR, ¹³C NMR and XPS techniques.

Staudenmaier Method

In 1898 staudenmaier improved Brodies' reaction by adding sulfuric acid to increase the acidity of mixture and KClO₃ over the reaction process[22]. Detailed procedure is briefed in Fig-4. In this method KClO₃ acts as an oxidant[23]with fuming HNO₃ as reaction media and the product contains Carbon-to-oxygen ratio as 1.17, Raman Spectral Ib/Icratio as 0.89, charge transfer resistance (Rcr)^a (k Ω) 1.74. In 2013 Sheshmani et al[24]modified Staudenmaier method for preparation of Graphene Oxide where they examined the effect of HNO₃/H₂SO₄amount on interlayer spacing and its comparison. In this method oxidation of Graphite with Nitric acid and sulfuric acid in 1:3 ratio for 4 days was performed which resulted in a slightly expanded interlayer distance to 9.13A $^{\circ}$.

Electrochemical Exfoliation methods for GO preparation

Electrochemical exfoliation methods for the preparation of Graphene oxide (GO) involves the use of electrochemical techniques to exfoliate graphite and introduce oxygen containing functional groups. In the initial steps for exfoliation of graphite there is intercalation of ions which is followed by expansion induced by gaseous products. The use of high anodic potentials creates oxidized Graphene (Graphene Oxide) which is rich in oxygen functional groups and structural defects.

Direct electrochemical exfoliation of Graphite for GO

In this method graphite is directly exfoliated in electrolyte medium for the production of high quality of Graphene oxide. Most commonly used electrolyte for this method are as sulfuric acid[25], Poly(styrenesulfonate) (PSS)[26]. Su et al[25]synthesized graphene oxide using electrochemical exfoliation method of graphite in sulfuric acid and obtained high quality graphene oxide. In this work natural graphite flakes were employed as an electrode and a Pt wire used as grounded electrode in H2SO4. The obtained graphene sheets exhibited ultra transparency (\sim 96% transmittance) and a resistance of <1K Ω /sq.

Ionic Liquid-assisted electrochemical exfoliation of Graphite

In this method ionic liquids are used in environmentally friendly electrolytes. In a study by Jamaluddin et al[27]produced surfactant-assissted exfoliated graphene oxide (sEGO) with 1-butyl-3-methyl-imidazolium (BMIM) and surfactant ionic liquids in presence of triple chain surfactant TC14 (sodium 1,4-bis(neopentyloxy)-3-(neopentylcarbonyl)-1,4-dioxobutane-2-sulfonate) where TC14 is substituted with BMIM generating surfactant ionic liquids to produce BMIM-TC14sEGO/NFC composites.





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Aqueous Electrochemical exfoliation of Graphite for GO preparation

In this exfoliation aqueous medium electrolyte is used. As graphene prepared by Anurag et al[28] prepared Graphene in an electrolytic bath which contains 0.1 M Ferrous sulfate (FeSO₄), two graphite rods are used at 3 cm distance and consistent potential of 15-20 V was applied across the setup and the resulting exfoliated graphene exhibited at 267 nm which is recognised to π - π transition of C-O bonds. Qingtao et al[29] used aqueous medium electrolyte method for preparation of exfoliated graphene oxide films where graphite is treated with concentrated sulfuric acid and hydrogen peroxide at room temperature then this exfoliated graphite without removing H₂SO₄ contained in a container which have holes for electrochemical oxidization, purpose of holes in walls is to confine graphite particles in limited space which avoid shedding of graphite from the reacting region during the process. On characterization it was found that the reduced exfoliated graphene oxide have a high conductivity of 1022 s cm⁻¹.

Surfactant assisted exfoliation of graphite

Two types of surfactant can be used in this exfoliation technique anionic surfactant and cationic surfactant both were tried and tested by Yang et al[30] where graphite is mildly oxidized using improved hummers method[31] and exfoliation is performed using HCDDACI and HCSL with different concentrations and they concluded that anionic surfactant was more favourable than cationic surfactants. Surfactants have a great deal on the extent of exfoliation, surfactants affect the exfoliation of mildly oxidized graphite through intercalating into the lamellas to weaken or strengthen the interlayer force.

Microwave assisted exfoliation method for preparation of Graphene oxide

Preparation of graphene via Hummers method, Brodies method or Staudenmaier method is a long time taking processes which involves highly toxic materials. Microwave assisted method is really time saving method for preparation of Graphene Oxide, GO was manufactured in 2015 by Viana et al.[32] via this process, which involved magnetically mixing natural graphite, 70% nitric acid, and KMnO4 in a glass flask for 10 minutes at ambient temperature. The graphite:HNO3:KMnO4 weight ratio was 1:2:1. Expanded graphite (EG) was made by transferring the acid dispersion to a crucible, heating it to 110°C for three hours, and then rinsing the resulting powder with deionized water until it became neutral. 2.2 ml of H2SO4 and 1.0 g of KMnO4 are added to 1.0 g of EG for oxidation and exfoliation. The mixture is magnetically agitated for 30 minutes to create a green dispersion, and then deionized water and the dispersion are sonicated for two hours, turning the brown color of the combination into a bright yellow tint. The mixture is then rinsed with 10% HCl and deionized water until it becomes neutral, resulting in a brown powder which is precipitated GO.Wang et al[33]Weighed and transferred 0.125g graphite in a 500 ml volumetric flask and 10 ml H2SO4 was added followed by the addition of 0.923g KMnO4. Stirred the mixture for 5 min then placed in Microwave. The microwave for 3 cycle and 30 sec each cycle for a total reaction time of 150 sec. After each cycle, keep the volumetric flask in ice bath for rapid cooling. Now 15 ml 30% H2O2 added drop wise and continue the reaction for 2 min with stirring, then centrifuge the solution for 10 min at 13000 RPM. Discard the supernatant and wash the solid with 5% HCl solution followed with ultrapure water as long as the pH of the solution is neutralized. The product has been dispersed in 100 ml water with stirring for 5 min. and used the ultrasonic bath to exfoliate the suspension for half an hour. The solution is centrifuged for 15 min at 4500 rpm and finally dry the solid product overnight at room temperature.

CONCLUSION

After reviewing all the procedures and chemicals used in all processes we conclude that all procedures have their own benefits for the production of GO and in this time of global warming and pollution fast and facile green methods should be used for the production of GO on commercial scale as the method and procedure discussed in this review gives 100% yield as claimed by the authors which takes very small time as compared to well established previous methods like hummers method and brodies methods.





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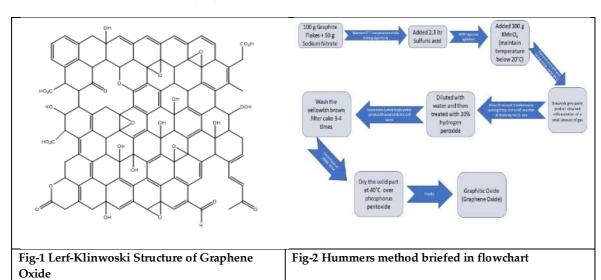
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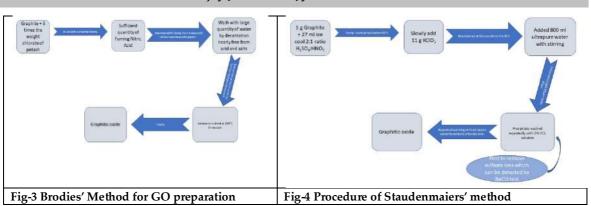
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REVIEW ARTICLE

A Review of House Dust Mites (HDMs) induced allergy, its Diagnosis and Treatment

Asha Poonia, Nisha Dhull, Sharda Kalra, Parvati Sharma, Vikash, Amit Kumar* & Kapish Samota

Faculty of Life Sciences, Chaudhary Bansi Lal University Bhiwani, Haryana, India.

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*Address for Correspondence **Amit Kumar**

Faculty of Life Sciences, Chaudhary Bansi Lal University Bhiwani, Haryana, India.

Email: amitmehra455@gmail.com



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ABSTRACT

Approximately 1-2% population of worldwide has allergic sensitization towards HDMs. Symptoms of HDMs allergens are usually respiratory related problems. Both innate and adaptive immune response activates against HDM allergens. Avoidance of allergens, pharmacotherapy treatment, Allergen immunotherapy (AIT) and probiotics are mainly used for treatment against respiratory allergies. Accurate diagnosis and designing of specific immunotherapeutic vaccine are mainly based on proper identification of allergens. This review describes the HDM characters, symptoms, diagnosis and treatments strategies with mechanism.

Keywords: House Dust Mites, pharmacotherapy, Allergen immunotherapy, probiotics, symptoms, desensitization, acaricides, India.

INTRODUCTION

House Dust Mites (HDM) are 0.2-0.4 mm in size tiny creatures. They are members of the suborder Astigmata in the class Arachnids in the kingdom Animalia. These pale creatures have a translucent body that ranges in size from 250 to 350 µm. They are a major indoor allergen source (Wong et al., 2011; Dey et al., 2019). They have a close relationship with people; high concentrations of HDM allergens are especially common in homes, carpets, and sleeping quarters. HDMs are primarily found in high-furniture areas like beds, chairs, and long-fiber carpets. They also feed on skin scales from human and pet activity, dander, and other materials such as household dust and plant particles (Tilak et al., 1994). In contrast to pollen, mite allergens specifically cause sleep disruptions, which lead to daytime lethargy and diminished bodily function (Brehler, 2023). Dermatophagoidespteronyssinus (Dp) and Dermatophagoidesfarinae (Df), two prominent pyroglyphid HDM species, are closely linked to allergic disorders (Dey et al., 2019). Due to their enzymatic activity, these mites take advantage of compromised epidermal skin





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barriers in atopic dermatitis (AD) sufferers to get access to immune cells. The symptoms of AD are made worse by the HDM allergens, which cause both delayed and acute allergic reactions (Bussmann et al., 2006). While HDMs are most prevalent during the rainy season, their greatest influence is seen in the drier and cooler months. It has been determined that the ideal temperature range for their development is between 25 and 26°C (AICPAHH, 2000; Colloff, 2009; Mumcuoğlu and Özkan, 2020). Gaining an understanding of the complex interactions between HDMs and human habitats, as well as their propensity to cause allergies, is essential to understanding the dynamics of indoor allergen exposure and its effects on human health. With a short life cycle marked by fast reproductive turnover and a 7–10week life expectancy, house dust mites (HDMs) are made up of several primary species, including Dermatophagoidespteronyssius, Dermatophagoidesfarinae, and Blomia tropicalis.

Female HDMs can lay 40–80 eggs, which starts a life cycle that consists of five stages: mating, laying one or two eggs a day, and developing through larval stages (protonymphs, tritonymphs) until they become adults, either male or female (Yang and Zhu, 2017). This entire life cycle takes about a month to complete. 1-2 percent of people worldwide exhibit allergic sensitivity to HDMs, which primarily presents as respiratory symptoms. Asthma and rhinitis are common symptoms that are frequently ignored until they worsen. Sneezing, itching, watery eyes, wheezing, Headaches, exhaustion, and depression, Headaches, exhaustion, and inflammation around the neck area are among the symptoms. (Lyon, 1991). Hypersensitive people who inhale HDM allergens may experience acute bronchial asthma attacks, characterized by wheezing, dyspnea, and, in extreme situations, death (Sharma et al., 2011). Effective therapy and prevention efforts require an understanding of the prevalence of allergic sensitization to HDMs as well as the respiratory symptoms that are associated with it.

House Dust Mite Allergens: Optimizing Human Immune Responses"

Both innate and delayed type's immune response activates against HDM allergens (Bussmann et al., 2006). HDM allergens mainly affects the respiratory epithelial cells and mainly follows two different immunological pathways for allergenic effects: first, through the CD4 and TH2 cells that activate IgE dependent allergy responses and they also activate mast cells independently of IgE and second, through activating the innate immune system by proteases, immunological epitopes, structural polysaccharides chitin and compounds and ligands derived from mite-associated sources. Strong allergens stimulate the immune system's innate and adaptive components(Jacquet et al., 2013; Burks et al., 2013).

House Dust Mite (HDM) Diagnosis and treatment

Allergies pose difficulties in general care because they are frequently neglected until symptoms become severe. The diagnosis of HDM allergy has historically depended on a patient's history of HDM exposure, in addition to proof of HDM sensitization discovered by specific IgE testing and skin prick tests. To improve diagnosis accuracy, nasal challenge tests may be used (Pfaar et al., 2014; Calderon et al., 2015a). However, obtaining an accurate diagnosis demands increased knowledge at the primary care level. When clinical signs such as nasal obstruction, rhinorrhea, sneezing, and pruritus, particularly redness, are present, HDM allergy is diagnosed (Zhang et al., 2018). Improving diagnostic techniques and raising primary-level awareness are critical for addressing the complexities of HDM allergy diagnosis (Table 1). Currently, there are four primary therapeutic approaches available to treat respiratory allergic illnesses caused by HDM: (1) avoidance of allergens; (2) pharmacological intervention; (3) allergen immunotherapy (AIT); and (4) use of probiotics to combat respiratory allergies. Avoiding allergens and adopting AIT are only possible once the particular allergy has been identified.

Avoidance of allergens

Immunostimulatory proteins found in dust mite faeces, which are mostly found in most dust reservoirs in our homes—including mattresses, couches, carpets, and clothing—are the primary cause of allergies (Zuiani and Custovic 2020). The main strategies for managing HDM allergy include Allergen Avoidance, which include covering beds and carpets, keeping ventilation adequate, cleaning the house, using acaricides, and avoiding humid and hot environments (Mumcuoğluand Özkan ,2020).





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Pharmacological intervention

When used in pharmacotherapy, allergen extracts from allergic patients increase their standard of life and give longterm symptom alleviation during subsequent natural exposure. When treating individuals with IgE-mediated allergy disorders, it is quite successful. About 85% of patients were prescribed oral antihistamines and nasal steroids for the treatment of rhinitis and asthma. Other medications, such as oral steroids, oral LABA, oral corticoids, cromones, anti-leukotrienes, and theophylline, were rarely prescribed to rhinitis and asthma patient(Trebuchon et al., 2014; Calderón et al., 2015b; Demoly et al., 2016). At present, allergen immunotherapy (AIT) completely replaces it. By giving certain allergens to patients, allergen immunotherapy (AIT) strengthens their immune systems and helps them recover from allergic disorders. Co-exposure to substances can result in complex physiopathology of allergic disease due to a variety of factors, including riskpathway, allergen/antigen structural features, dose and exposure duration of allergens, genetic susceptibility and stimulators of the innate immune response like bacterial infections (Akdis and Akdis, 2014). AIT begins with identifying a patient's particular allergen, which can be a very challenging task at times due to the cross-reactivity among allergens (Zhang et al., 2018). Allergen immunotherapy (AIT) is one measure of the condition's severity. The two main categories of AIT are SLIT and SCIT. The primary goal of subcutaneous immunotherapy (SCIT) and sublingual immunotherapy (SLIT) is the delivery of immunotherapy, namely allergen-specific immunotherapy or allergy-specific immunotherapy (Burks et al., 2013). For the majority of outcomes, SCIT and SLIT were more beneficial than placebo. Medication and symptom scores improved more with SCIT than with SLIT, and there were no changes in quality of life. One effective alternative for treating allergic rhinitis, including HDM-triggered AR, is allergen-specific immunotherapy (AIT). According to Eifan et al. (2013) and Fujita et al. (2012), allergen tolerance Th2 responses are suppressed when AIT activates Treg cells. By modifying the immune response to AIT, adjuvants also improve efficacy and safety.

The allergen immunotherapy AIT Mechanism

Events that occur at the cellular and molecular level during AIT can be divided into three categories.

- 1. Tregulatory (Treg) and B regulatory (Breg cells) promote switching of B cell class
- 2. Suppression of late-phase effector cells (Eosinophils & T-cells) and
- 3. Rapid desensitisation of mast cells and basophils

Treg and Breg cells promote B cell class switching

The effectiveness of AIT depends on the activation of Breg and Tregcells . Originating in the thymus, teg cells regulate both tissue-specific and systemic autoimmune. IL-10 is essential for effective immunotherapy since it is linked to T-cell tolerance in allergic disorders. Breg cells have potent regulatory and suppressor capabilities, especially IL-10-secreting B cells The release of IL-10 after AIT and spontaneous allergen exposure is caused by antigen-specific IL-10-secreting regulatory T (TR1) cells and IL-10-secreting regulatory B (BR1) cells(Van et al., 2013). T and B regulatory cells stimulate B cell class switching, which influences IgE production. Serum-specific IgE levels are momentarily elevated by AIT before declining with continued treatment. Improved clinical outcomes are correlated with elevated Specific IgG, particularly IgG4. By preventing the release of inflammatory mediators that are induced by allergens, blocking antibodies also hinders the formation of memory IgE with increased allergen exposure and reduces the ability of IgE to deliver allergens to T cells (Akdis and Akdis, 2014).

Suppression of late-phase effector cells (Eosinophils & T-cells)

AIT also inhibits late-phase effector cells linked to late-phase responses (LPR), including T cells and eosinophils. According to Van and Stevens (1989) and Akdis and Akdis (2014), AIT increases the concentration of allergen needed to cause instant reactions, or LPRs, which reflects underlying mucosal inflammation and correlates with clinical improvement. Reduced responses to AIT indicate that the LPR interaction is essential to its therapeutic efficacy. As demonstrated by reduced responses to non-stimulatory stimuli and improvements in bronchial, nasal, and conjunctival hyper-reactivity following AIT, the interaction between AIT and the LPR is essential to its therapeutic success (Akdis&Akdis, 2014).





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Rapid desensitisation of mast cells and basophils

As part of allergen immunotherapy, mast cell and basophil desensitisation quickly is essential (AIT). After the first treatment with allergen supplements themast cells and basophil activity has significantly decreased, which makes them immune to environmental proteins—even when particular IgE is present (Akdis and Akdis, 2014). Despite high levels of specific IgE in treated participants, systemic anaphylaxis can occur early in the AIT process, especially after the initial injection of three-dimensional structure-intact allergens due to a decrease in sensitivity in mast cells and basophils. The degree of mediator release from these cells is changed by successful hypo-sensitization, which affects the activation thresholds that follow (Boonpiyathad et al., 2019).

Application of probiotics for respiratory allergies

The composition of the immune system is determined by the microflora in the gut, and probiotic-based microflora enrichment can be very important in preventing allergic reactions (Zuccotti et al., 2015; Ismail et al., 2013). The microbiota in the stomach suppresses the synthesis of T helper-1 (T1) cytokines and modifies their reaction. Additionally, the microbial fauna suppresses the T2 response and induces T regulatory cells (Smith-Norowitz and Bluth, 2016). Probiotics boost the production of IgA locally, which impacts mucosal defence. In order for adaptive T cell immunity to mature, they also promote the production of IL-17 (Isailovic et al., 2015). Thus, probiotics are also crucial in preventing HDM allergic reactions, based on recent research (Fassio and Guagnini, 2018). Heat-killed Lactobacillus plantarum Probiotics such as Lactobacillus plantarum, Lactobacillus paracasei, Lactobacillus rhamnosus, Bifidobacterium lactis BL-04, Lactobacillus acidophilus NCFM, and Clostridium butyricum are being utilised to treat allergic disorders.

House dust mite sensitization detection Skin prick test

Due to its high sensitivity, the skin prick test with mite extracts is the best approach for determining house dust mite sensitization. The test outcome is dependent on the quality of the allergen extract, which must include all relevant components in sufficient quantities. Skin testing has traditionally been the gold standard for allergy diagnosis, offering quick results for discussion with patients and eliminating the need for follow-up meetings or laboratory reports (Heinzerling et al., 2013; Gureczny et al., 2023; Brehler, 2023).

In vitro tests (detection of specific IgE antibodies)

In vitro assays entail the use of common commercial techniques to identify certain IgE antibodies. For reliable findings, the allergen extract's purity is essential. It is possible to identify specific IgE in nasal secretions in cases of suspected local allergic rhinitis, though this has not been confirmed. Dermatophagoidespteronyssinus and farinae extracts are available for in vitro testing along with key allergens such as Der p 1, Der p 2, Der p 23, and Der p 10 (Weghofer et al., 2013; Haxel et al., 2017).

Provocation test

An allergy diagnosis must be confirmed by a nasal or conjunctival provocation test in circumstances when the patient's history is unclear. Subjective symptom scores and objective measures made with methods such as peak nasal inspiratory flow determination, acoustic rhinometry, anterior rhinomanometry, and 4-phase rhinomanometry are included in the evaluation (Haxel et al., 2017; Brehler et al., 2023).

Global and Indian Perspectives on Current Research and Future Prospects in Allergy Studies

Rigidity of the respiratory system affects almost one-third of the population in India, and its detrimental effects on quality of life have a major socioeconomic cost. According to epidemiological data, almost 25% of people in India have sensitivity to several allergies, such as rhinitis and asthma. A comprehensive approach is required to manage and comprehend the increasing prevalence of allergies, including eczema, angioedema, food and insect allergies, and anaphylaxis. Due to a lack of sophisticated multiplexed assay systems, there are few reports on developments in molecular allergology; nevertheless, India is now placing a higher priority on the creation of instruments and





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approaches for improved allergy control. In order to understand immunological mechanisms and provide better treatments, molecular allergy research conducted in India focuses on the detection, identification, purification, and molecular characterization of individual allergen molecules, including recombinant allergens. Nevertheless, because there aren't enough qualified clinical allergists and well-equipped allergy centres, problems with applying allergen immunotherapy (AIT) still exist. To address regional differences in allergenicity, proper identification, purification, and molecular characterization of House Dust Mite (HDM) allergens are stressed. In comparison to Europe and the United States, Asia has a greater global burden of dust mite sensitization. All countries use different approaches, although diagnostic technologies like Immuno CAP, allergen microarrays, and skin prick testing are widely employed. As a treatment for IgE-mediated allergy illnesses, AIT is still widely used since it produces long-lasting immunological and clinical tolerance. Probiotics are being investigated as a potential substitute to lessen HDM symptoms, strengthen mucosal defences, and enhance allergy sufferers' quality of life. The best bacterial strains for reducing allergy symptoms are still being researched globally.

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Table 1. Clinical Cases of Allergy: Identification and Customised Immunotherapeutic Treatments"

Case No.	Scenario of Clinical Cases	References
1.	The patient in the first scenario had many lesions spread across her body, especially on her hands, feet, scalp, and trunk. These lesions are suggestive of severe refractory atopic dermatitis. Using a diagnostic technique, the condition is identified thorough evaluation. The process of treatment consists of a combination of medicinal approaches, including systemic steroids, antihistamines, immune suppressants, moisturizers, emollients, and antibiotics. Notably, sublingual immunotherapy (SLIT) is advised since it effectively reduces adverse effects in patients.	Yang et al., 2022; Huang et al., 2023; Brehler, 2023
2.	In the second scenario the symptoms include a red rash around the neck, watery eyes, coughing, sneezing, and itching, all of which are indicative of an infection that has been linked to house dust mites (HDMs). To effectively manage the condition, the diagnosis entails determining the causing factors. Treatment options include corticosteroids and moisturizer lotions, with an emphasis on inhaling dust mite allergens.	Yang et al., 2022; Brehler, 2023; Batard et al., 2023
3.	The third scenario is about bronchial asthma brought on by a lifetime of exposure to allergens related to HDM. Oral antihistamines and nasal steroids are part of the pharmacotherapy suggested to effectively control the illness and relieve symptoms.	Zuiani, C. and Custovic, 2020
4.	The patient in the fourth case exhibits lesions and a rash suggestive of a long-term HDM infection. The preferred course of treatment is Allergy Immunotherapy (AIT), which offers long-term advantages beyond the course of the treatment and the ability to change the course of the allergic disease naturally.	Zuiani, C. and Custovic, 2020; Yang et al., 2022; Brehler, 2023; Batard et al., 2023
5.	Swelling and redness are indicative of mild to moderate allergic reactions in the fifth case. The recommended course of action is called Specific Allergen Immunotherapy (SCIT), which is designed to provide the patient with focused relief from allergic rhinitis and asthma. The significance of accurate diagnosis and customised treatment plans in the management of various allergy diseases is highlighted by these organized interventions. (Sources: a) SCIT for mild to	Yang and Zhu, 2017; Zuiani, C. and Custovic, 2020; Yang et al., 2022; Brehler, 2023; Batard et al., 2023





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moderate HDM infection, b) Inhalation therapy for HDM infection, c) AIT for chronic HDM infection

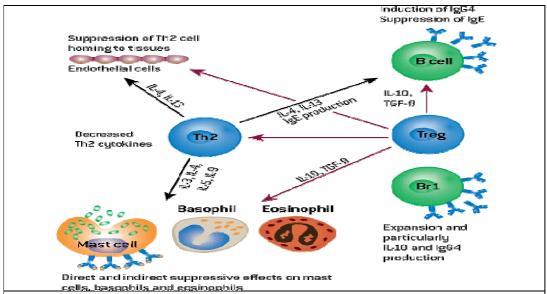


Fig.2. from (Akdis and Akdis, 2014).The complex regulation of allergic inflammation is carried out by Treg and Breg cells. The equilibrium between TH2/Treg cells is a critical factor in the development of allergic inflammation. Allergy-related disorders are impacted by TH2 immune response suppression by treg cells and associated cytokines. The regulatory effects of Treg cells are indicated by red arrows. These effects include reduction of IgE and induction of IgG4/IgA and B cell homing, as well as direct and indirect regulation of mast cells, basophils, and eosinophils and vascular endothelium. Breg cells also promote the generation of IgG4 and inhibit effector T cells.





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RESEARCH ARTICLE

A brief overview on Schiff Base Synthesis techniques

Gurdeep Sangwan, Jyoti Sharma*, Avinash Rani and Sonu Prasad

Department of Chemistry, MMEC, Maharishi Markandeshwar (Deemed to be) University, Mullana-133207 (Ambala) India.

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*Address for Correspondence Jyoti Sharma

Associate professor,

Department of Chemistry,

MMEC, Maharishi Markandeshwar (Deemed to be) University,

Mullana-133207 (Ambala) India. Email: jsharma117@gmail.com



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ABSTRACT

This comprehensive review delves into the synthesis of Schiff bases, pioneered by Hugo Schiff in the 19th century, elucidating their significance in organic chemistry. Schiff bases, characterized by the C=N group, result from the condensation of primary amines with active carbonyl compounds. The review explores classical and environment friendly approaches, dissecting methods such as condensation with Lewis acid, use of molecular sieves, and azeotropic distillation. Additionally, non-conventional methods like water-based synthesis, microwave-assisted synthesis, ultrasound-promoted synthesis, and grindstone chemistry are discussed, emphasizing advancements in reaction efficiency, selectivity, and sustainability.

Keywords: Schiff bases, organic synthesis, condensation, environment friendly approaches, nonconventional methods, reaction efficiency.

INTRODUCTION

In the 19th century, Hugo Schiff pioneered the synthesis of Schiff bases, also known as imines[1]. ASchiff base is defined as an aldehyde containing a C=N group (azomethine -N=CH- group) as shown in Figure 1[2]. These are synthesized by a condensation reaction of primary amines and an active carbonyl compound (aldehydes/ ketones). Water is the by- product of this reaction. The general formula of Schiffbase is RCH=NR1, where R & R1 represents alkyl/aryl substituents [3]. Compounds featuring anazomethine group exhibit basic properties owing to the presence of unshared electron pairs on the nitrogen atom and the electron-donating characteristic of the double bond. Schiff bases, a subset of compounds with an azomethine group, display relatively weaker basic properties when compared to their corresponding amines. This disparity arises from the alteration in hybridization during the formation of the imine structure. While the nitrogen atom in amines undergoes sp³ hybridization, the hybridization shifts to sp² in the





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imine structure. As a consequence of this increased s character in hybridization, there is a significant reduction in basicity. Schiff bases are selective toward metal ions and form complexes by transferring electrons from the active ends they contain to the metal. Schiff bases are known as a good nitrogendonorlig and (—CH=N—). During the formation of the coordination compound, one or more electron pairs are donated to the metal ion by these ligands. Schiff bases can form highly stable 4-, 5-, and 6-ring complexes if they donate more than one electron pair. For this, a second functional group with a displaceable hydrogen atom must be found as close as possible to the azomethine group. This group is preferably the hydroxyl group or any S atom Schiff base compounds play a crucial role in both pharmaceutical and chemical industries, showcasing a spectrum of biological activities such as antimicrobial, antidyslipidemic, antitubercular, antihelmintic, anticonvulsant, anti-inflammatory, antiviral, antioxidant, and more[4]. These compounds have been a focal point in organic synthesis for decades. The classical condensation reactions involving primary amines and aldehydes or ketones have paved the way for the synthesis is of these valuable compounds. However, as the field advances, researchers are exploring novel methodologies to enhance reaction efficiency, selectivity, and environmental sustainability. In this article we are going to explore conventional and non-convential approaches for the preparation of Schiff base as shown in Figure 2.

Different methods of Schiff base synthesis Conventional methods of Schiff base synthesis

Condensation in the presence of Lewis acid: The synthesis of the Schiff base is carried out by condensing amine with aldehyde in suitable solvent, such as ethanol at refluxing. There action is carried out in the presence of a catalyst: Lewis acid. The mechanism of the reaction is shown in figure 3[5]. This method is widely used in the synthesis of Schiff base. However, this method has a limitation of performing reaction at high temperature and longer duration. A Schiff base was synthesized through the reaction of an equi molar mixture of 4-amino benzene sulphon amide and substituted aromatic aldehydes or acetophenone. The individual reactants were dissolved in a minimal amount of ethanol, combined, and supplemented with 2 ml of glacial acetic acid. The resulting solution underwent reflux for 8 hours, followed by cooling to room temperature and pouring into ice-cold water. The solid precipitate obtained was collected via filtration, dried using an oven at 80 °C, re dissolved in ethanol for re crystallization, and subsequently dried to yield the final product[6].

Using molecular sieves/ azeotropic distillation for water removal

In Schiff base synthesis, molecular sieves play a crucial role by serving as desiccants to remove water from the reaction mixture. The presence of water can adversely affect the reaction equilibrium and lead to hydrolysis of the desired imine product. To mitigate this, the water can be removed by using Dean–Stark apparatus azeotropically. Also, molecular sieves, commonly of 3 Å or 4 Å pore size, are introduced into the reaction vessel along with the aldehyde or ketone and primary amine. As the condensation reaction progresses, water is produced, and the molecular sieves continuously adsorb and remove it, promoting the formation of the Schiff base. This ensures a more efficient and higher-yielding synthesis. Following the completion of the reaction, the molecular sieves are removed, and the Schiff base product can be isolated through standard workup procedures. Proper care, including pre-drying of molecular sieves and choosing the appropriate pore size, contributes to the compliance of the reaction. Overall, molecular sieves serve as effective tools in facilitating Schiff base synthesis by maintaining a dry reaction environment. This process is also time consuming and the molecular sieves need to be dried at high temperature to make them moisture free. Kazuo Taugchi and F.H. Westheim utilized molecular sieves for the synthesis of ketimines, as described in their work [7]. In their experimental procedure, around 40 grams of Linde 5Å molecular sieves were introduced into a mixture comprising 0.10 moles of ketone (Cyelooctanone, Cycloheptanone, Cyclohexanone, camphor, keto acids, and Acetophenone) and 0.12 moles of aromatic amine(Anilines and m-Toluidine) dissolved in either 40 milliliters (about 1.35 oz) of benzene or ether. There action duration spanned 1 to 30 hours, during which the progress was monitored through infrared (IR)and nuclear magnetic resonance (NMR) spectroscopy to ensure the absence of ketones. The isolation process encompassed filtration, removal of the solvent, and subsequent purification utilizing vacuum distillation or crystallization techniques.





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Non-conventional methods for Schiffbase synthesis Water as green solvent

This includes the use of water as a solvent and environmentally benign catalysts to minimize the environmental impact. Using water as solvent is also safe because it is non- toxic and non-flammable. Rao and colleagues [8] showcased an efficient method for synthesizing Schiff bases in an aqueous environment, yielding superior results compared to conventional approaches. Notably, this technique obviates the need for acid catalysts. The Schiff bases produced exhibited high yields and could be easily separated through filtration. It is noteworthy that water efficiently facilitated the reactions. This innovative method significantly accelerated the reaction rate, achieving a nearly 300-fold increase. The procedure involved adding salicylaldehyde to a solution of 1,2-diaminobenzene in 10 milliliters of water, followed by stirring the mixture for 10 minutes at 25 °C. The resulting yield was an impressive 95%, surpassing the 65% yield obtained through the traditional method.

Microwave-Assisted Synthesis

Microwave technology utilizes electromagnetic waves generated by a magnetron, typically at 2.45 GHz to avoid interference. During microwave irradiation, polarized molecules align with the electric field, inducing molecular motion and generating heat (dipolar polarization). Ionic conduction, involving charged compounds, also contributes to heat generation through oscillating trajectories. Temperature in the reaction mixture can peak at 250-300 °C, facilitating faster reactions based on the Arrhenius equation. Synthesis instruments often use sealed tubes for pressure buildup. Solvent choice is critical for microwave absorption and heat generation. In organo metallic chemistry, metals or metallic salts ensure microwave absorption, promoting reactions through localized temperature increases (hot spot theory)[9]. It presents advantages such as expedited reaction rates, higher yields, energy efficiency, and reduced reaction durations. By directly heating the reaction mixture, microwave irradiation facilitates rapid and efficient warming, leading to accelerated reactions and improved product quality. Microwaves have the capability to directly penetrate the reactants, inducing heat through molecular rotation and frictional heating. Mousumi Chakraborty et al[10] used microwave irradiation method, in which, reactant mixtures were subjected to microwave power at 300 W from 30 sec to 2 min and temperature was maintained at 80°C. The progress of the reaction at different time interval was monitored by TLC. On completion of reaction, the yellow coloured amorphous product 1 was separated, filtered, dried and re crystallized from methanol. There action occurred within very short time (30sec-2min).

[1]

In another microwave-assisted synthesis, performed by Bhusnureet al, a mixture of 3,5-dichlorobenzaldehyde (0.001 mol), 3-amino-6-bromo-2-phenylquinazoline-4(3H) one (0.001 mol), and two drops of ß-ethoxyethanol as a wetting reagent was prepared in a beaker. The reaction mixture underwent irradiation within a modified MC767W (Electrolux) system at 200 W for approximately 3minutes, with intermittent cooling intervals, Reaction was monitored by TLC. After the 3-minutereaction period, the flask was cooled in ice water, diluted with ice-cold water, and the resulting Schiff bases was filtered, dried, and re crystallized from ethyl acetate. Using the same procedure, other compounds were synthesized within 3-5 minutes (180-300 seconds). Optimization involved varying the reaction time and microwave power from 180W to 600W. The conversion and yield was found to be better than that with the conventional method of synthesis[11].





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Ultrasound-Promoted Synthesis

Sono-chemistry refers to the use of ultrasonic vibrations to enhance chemical reactivity. Ultrasound irradiation is a green technique used to speed up chemical processes, particularly inorganic synthesis. Ultrasonic waves, with frequencies above 20KHz, are particularly useful for chemical processes. As sound waves travel through a liquid medium, molecules oscillate due to cycles of rarefaction and compression. Under certain conditions, bubbles can emerge, leading to high local temperatures and pressures. Ultrasonic cavitation occurs when low pressure is applied to a liquid medium, causing cavities or bubbles to form and collapse. This technique is commonly used in laboratories using an ultrasonic bath or probe[12]. This technology is easier to handle and more convenient than older methods. With ultrasonic irradiation, many chemical processes can be carried out in milder conditions and with shorter reaction periods, yielding greater yields [13]. When compared to traditional heating, this can be seen as a processing help in terms of waste reduction and energy conservation. Hadi Kargar et al[14] have successfully synthesized and characterized two Schiff base compounds derived from 3-ethoxysalicylaldehyde and 2aminopyridine derivatives under ultrasound irradiation. Compound 2 was prepared by placing a 1:1 M combination of 3-hydroxy-2-aminopyridine and 3-ethoxysalicylaldehyde in ethanol in an ultrasonic bath set at room temperature for five minutes. TLC was used to monitor the reaction's development. Subsequently, the mixture was exposed to air to gradually evaporate over a few days, allowing the products to settle. After characterization by NMR spectroscopic techniques and elemental analyses, the crystal structures of compounds were confirmed by single crystal X-ray analysis.

[2]

A Schiff base was synthesized by Ahmad Aziz and co-workers by condensing piperonal (15 mmol)and anthranilic acid (15 mmol) with acetic acid as a catalyst in methanol. The mixture was sonicated at 37KHz for 10 minutes at 25 °C in an ultrasonic bath, and the reaction progress was monitored using TLC. After completion, the reaction mixture was concentrated, cooled, and filtered. The resulting precipitates were washed with methanol, resulting in a yellow product. The structure of the Schiff base was confirmed through analysis of physical and spectroscopic data[15].

Grinding method

Many exothermic reactions can achieve high yields simply by grinding solid reactants together using a mortar and pestle, a method referred to as "grindstone chemistry." This technique, categorized under "green chemistry," involves initiating reactions through the act of grinding, where minimal energy transfer occurs through friction. Grindstone chemistry not only proves energy-efficient but also exhibits heightened reactivity and produces fewer waste products. These reactions are straight forward to manage, contribute to pollution reduction, are relatively cost-effective, and are considered more economical and environmentally friendly procedures in the field of chemistry. Solid-state reactions, facilitated by grindstone chemistry, demonstrate enhanced efficiency and selectivity compared to solution reactions, owing to the tightly arranged and regular molecular structure in crystals[16]. The solid-state synthesis of several types of benzylidene aniline derivatives without going through liquid phases was described by Schmeyers et al [17]. Equi molar mixtures of pure aniline derivative and aldehyde derivative were ground together in a mortar for two hours at room temperature to carry out the solid-solid reactions. The completion of the reaction was checked by IR spectroscopy in KBr. The water produced in there action was removed at 80°C under vacuum.





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Chemical analysis was carried out by IR and NMR spectroscopy which gave the expected peaks and signals. In contrast to (acid catalyzed)imine synthesis in solution, these solid-solid condensations occur without the production of waste.

CONCLUSION

In summary, this comprehensive review explores the evolution of Schiff base synthesis methodologies from classical to innovative approaches. While classical methods have proven effective, the emerging techniques, including microwave-assisted and ultrasound- promoted synthesis, water as a green solvent, and grindstone chemistry, offer advantages such as expedited reactions, higher yields, and environment friendly procedures. The dynamic pursuit of sustainability, efficiency, and selectivity in Schiff base synthesis reflects the ongoing commitment of researchers to advance the field of organic and inorganic synthesis.

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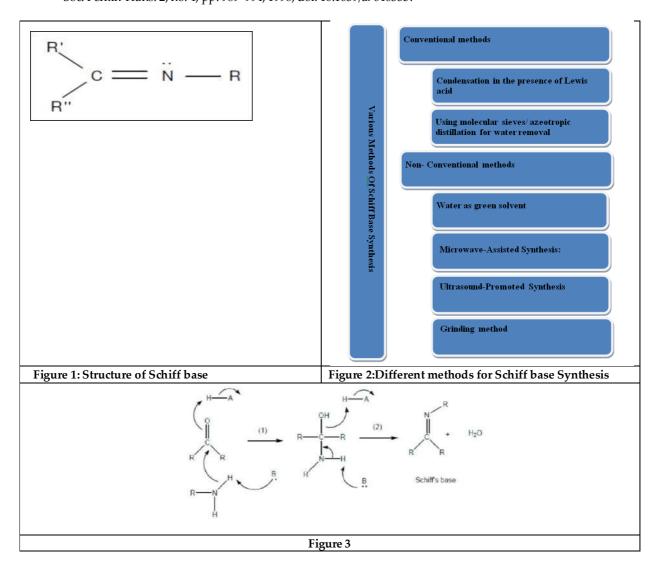
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RESEARCH ARTICLE

Chemistry of Iridium(III)-3-hydroxy-2-tolyl-4H-chromen-4-one: The Spectrophotometric Analysis, Radical Scavenging and **Computational Studies**

Khushboo Devi, Nivedita Agnihotri*, Gulshan Kumar

Department of Chemistry, Maharishi Markandeshwar (Deemed to be University), Mullana, Ambala-133207, India.

Received: 30 Dec 2023 Revised: 09 Jan 2024 Accepted: 12 Jan 2024

*Address for Correspondence Nivedita Agnihotri

Department of Chemistry,

Maharishi Markandeshwar (Deemed to be University),

Mullana, Ambala-133207, India.

Email: niveditachem@mmumullana.org;



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ABSTRACT

3-Hydroxy-2-tolyl-4H-chromen-4-one (HToC)has been utilized successfully for the microgram level analysis of iridium (III) employing a spectrophotometric determination technique. Iridium (III) instantaneously reacts with HToC in phosphoric acid medium to form a stable light yellow-coloured non-extractable complex. The Ir(III)-HToC complex evinces a maximum absorbance at 423nm when studied against a reagent blank prepared analogously and coheres to linearity up to 1.8 µg Ir (III) ml-1. The accuracy and reproducibility of the method are confirmed by the corresponding values of relative standard deviation (%RSD) and the coefficient of variance as 0.0146% and 0.00146, respectively, at 423 nm. The stoichiometric constitution of the developed complex, as validated by the studied Mole Ratio and Job's continuous variation methods is 1:3 [Ir(III):HToC], indicating hence the octahedral geometry of the formed species. Study has been done for interference by various cations, anions and complexing agents to analyse selectivity of the method. It is observed that the majority of the ions/complexing agents do not interfere with the complex determination. Besides, electronic properties of the studied complex have been affirmed by subjecting it to the computational studies. The practicability of the procedure has been assessed by conducting analysis of numerous synthetic samples. Finally, the studied complex is exposed to the estimation of antioxidant capacity by employing DPPH radical scavenging method. Antioxidant capacity results propounded the better radical scavenging capacity of the developed complex than that of the ligand alone.

Keywords: Iridium(III), spectrophotometric analysis, radical scavenging activity, computational studies, QTAIM, TDDFT





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INTRODUCTION

Organic and inorganic compounds including transition metals especially the platinum group metals (PGMs)are qualitatively and quantitatively determined by using numerous analytical methods, such as volumetric titrations [1], atomic absorption spectrophotometry (AAS) [2], neutron activation analysis (NAA) [3], UV-VIS spectrophotometry [4], ion exchange chromatography [5], inductive coupled plasma optical emission spectroscopy (ICP-OES) [6], X-ray fluorescence spectrometry[7] and voltammetry [8]. Among the several methods for determining the metals in traces, UV-VIS spectrophotometric methods ofdetermination are the most generally employed due to less apparatus and methodology requirements. Spectrophotometric approaches are regarded relatively convenient for micro level determination and are widely utilized for the same reason. The elucidation of results in UV-VIS spectrophotometer can be operated to analyze the chemical elements, structure and composition of inorganic complexes[4]. The method is based on the basic principle of measurement of intensity of ultraviolet and visible light absorbed by the sample as a function of wavelength. Samples absorbs ultraviolet radiation at a wavelength of 180-380 nm or visible at a wavelength of 380-780 nm. The results are estimated in the form of transmittance or absorbance or UV-VIS spectrum of the complexes prepared as clear solutions. All PGMs including ruthenium, rhodium, palladium, osmium, iridium and platinum crop up naturally in the Earth's crust at very low concentrations [9]. Iridium, one of the PGMs is known as a very hard, brittle and silvery-white metal. After osmium it is considered the second-densest (density 22.56 gcm⁻³) naturally occurring metal[10]. The metal was discovered in 1803 by Smithson Tennant.

Its chemical name originated from the Greek goddess, Iris, an epitome of the rainbow [11]. Iridium forms compounds in variable oxidation states ranging from -3 to +9, the most common oxidation states being +1, +3 and +4. Iridium has extensive coordination chemistry. Iridium in its complexes is always low-spin. Octahedral complexes of Ir(III) and Ir(IV) are generally known[12]. Iridium and its complexes find extensive applications in various fields such as catalysis, medical imaging, electrodes for producing chlorine and other aggressive products, OLED, crucibles and ignition tips for spark plugs [13-15]. Radioisotope, iridium-192, is one of the most important source of energy used in industrial y-radiography for non-destructive testing of metals and in the treatment of cancer using brachytherapy, a form of radiotherapy[16]. Iridium along with otherplatinum group metals play an important role in analytical chemistry producing a variety of complexes with a number of ligands carrying variable donor groups and studied by the spectrophotometric techniques. Coloured complexes of iridium are reported with hydrazine carbothioamide[17], p-methylphenylthiourea [18] and Schiff bases [19] but suffering with very poor sensitivity. However, 4H-1-benzopyrans were observed to show significant improvement when used as ligands over to the others[20,21]. For the same reason, in the present study, another benzopyran derivative, 3-hydroxy-2-tolyl-4Hchromen-4-one (HToC; figure1) has been employed as a noble reagent for trace determination of iridium in its trivalent oxidation state to give a very enhanced value of sensitivity as compared to the other mentioned reagents[17-19]. It has been searched out that the transition metal complexes with organic especially chelating ligands helped in enhancing the antioxidant potential while used in medicinal and pharmaceutical industries [22]. Chelation exhibited by the metal complexes is the key for this improved activity over the traditionally used synthetic antioxidants. On the same basis, in the present research the newly prepared Ir(III)-HToC chelate has been subjected to antioxidant potential and was found to betray the same [23].

Experimental section

MATERIAL AND METHODS FOR SPECTROPHOTOMETRIC ANALYSIS

Instrumentation and solutions

All the absorbance measurements were performed using UV-VIS spectrophotometer (double beam; EI-2375) with 1 cm identical quartz cuvettes. A 100 ml of the stock solution of iridium with a strength of 1 mg Ir(III) ml⁻¹and working solutions of the required concentration (100 and 10 μ g ml⁻¹) were prepared by the reported method [20]. High purity chemicals of AR grade and deionized water were used for all the experiments. For the regulation of the ambient





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acidity of the reaction medium, $0.5 \text{ M H}_3\text{PO}_4$ was prepared. Other required solutions were prepared to study the effects caused by foreign ions (cations, anions and complexing agents) from the respective sodium or potassium salts by dissolving in appropriate solvents.

Ligand solution

The chromogenic reagent, 3-hydroxy-2-tolyl-4H-chromen-4-one (HToC), carrying the carbonyl group and the hydroxyl group as the binding sites (figure 1) was synthesized by applying the reported methodology[24]. Itsfreshly prepared solution of strength 0.1% (w/v) was made by dissolving 0.05g of it in a small amount of ethanol, followed by dilution with the same solvent up to the mark in a 50 ml volumetric flask.

Spectrophotometric Determination

In order to achieve accurate and impeccable results, the spectrophotometric determination was carried out in the most optimized manner. By diluting the stock solution with appropriate amounts of double-deionized water, a working solution (10 μg ml⁻¹) of Ir(III) was prepared for analysis. For the process of complexation, 1 ml of the working solution of the metal ion in trivalent oxidationwas pipetted out in a 10 ml standard flask, 1 ml of the 0.1% chromogenic reagent solution in ethyl alcohol was added to develop complex in the presence of 0.035M H₃PO₄.After adding 1ml of ethanolic solution of 0.1% (w/v) HToC, a light yellow-coloured complex was formed instantaneously. The colour intensity of Iridium(III)-HToC complex was measured at 423nm against a comparable reagent blank solution. The precise amount of metal ion was calculated using the calibration curve constructed between absorbance and varying iridium(III) concentrations.

Antioxidant activity

The antioxidant capacity of Ir(III)-HToC was mannered by Radical Scavenging Activity (RSA) towards 2,2-diphenyl-1-picrylhydrazyl analysis (DPPH) and the outcomes were observed by engaging a UV-VIS spectrophotometer. DPPH radical scavenging is the well known procedure to the evaluation of the antioxidant property of chemical compounds. The DPPH (methanolic solution) gives deep purple colour contributing an acute absorption band at 517 nm; where colour change of solution is indication of the reaction with the substance that has antioxidant capacity. Stock solution of 1 mg ml⁻¹ of Ir(III)-HToC was prepared by dissolving 5 mg of it in 5 ml methanol. However, the stock solution of DPPH of strength 1.0 mM and the solution acting as a negative control were prepared similarly as reported [21,25]. A number of range of dilutions (500, 250, 125, 62.5 and 31.25 µg ml⁻¹) of complex were prepared from the stock solution of Ir (III)-HToC. To retain the RSA of the resultant complex and the ligand (HToC), lower dilutions of HToC of the same concentrations as those of Ir (III)-HToC complex were prepared consequently. This was accompanied by incubating the solutions at 37°C for 30 min in dark. The positive control, gallic acid of same concentrations were prepared simultaneously. Ultimately, the absorbance of each solution was spectrophotometrically determined at 517 nm against the prepared negative control, and the % radical scavenging activity was calculated as per the equation: % RSA = [(Anegativecontrol-Asample)/ Anegative control ×100, where Anegative control denotes the absorbance of negative control and Asample denotes absorbance of the antioxidant compounds (gallic acid, Ir (III)-HToC, HToC) [22-25].

Software details

The reagent HToC and Ir(III)-HToC complex were optimized at ground state (S_0) using density functional theory (DFT) at the B3LYP/ LANL2DZ level[26]. The absence of imaginary vibration frequencies analyzed the stability of molecular structures. The excitation energies were calculated using time-dependent DFT (TDDFT) methods. All DFT and TDDFT calculations were carried out using Gaussian 16 [27]. Quantum chemical parameters were utilized to investigate the electronic properties of the complex's most stable conformation. This was done to better understand the studied complex's chemistry. The electronic properties of HToC and Ir(III)-HToC complex, such as the ionization energy (I), electron affinity (A), electronegativity (χ), chemical potential (μ), absolute hardness (η), global softness (σ), and electrophillicity index (ω), were determined using the band gap energy ($E_{gap} = E_{HOMO} - E_{LUMO}$) and the HOMO/LUMO energies. Frontier molecular orbitals are used to determine these properties as main orbitals involved





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in chemical stability. A molecule's reactivity can be explained by considering its chemical hardness, which is directly related to the HOMO–LUMO energy gap. A molecule with a greater E_{gap} or chemical hardness is less reactive [28-30]. The application of theoretical studies on the complex was further expanded by employing QTAIM analysis[31].

RESULTS AND DISCUSSION

A light yellow stable complex (>2 days) ofIr(III)-HToC was obtained aqueous phase upon interaction with the chromogenic reagent, HToC from an acidic medium provided by phosphoric acid. The complex was non extractable in any of the solvents, polar or nonpolar, when subjected to its extraction. The spectrophotometric determination procedures, computational studies and bioanalytical analysis of the complex are mentioned in succeeding discussion.

Spectrophotometric analysis

Absorption spectrum

Under aqueous conditions, the absorption curve of Ir(III)-HToC complex against blank had a paramount and consistent colour intensity in the range 420–425 nm in phosphoric acid medium. The spectrum of reagent blank when examined similarly against pure water indicated a minimal absorbance in the same region (figure 2). Thus, 423 nm wavelength was fixed for absorbance in case of all further parameters and against reagent blank for the analysis of Ir (III).

Effect of medium and acidity

Of the numerous acidic (HNO₃, H₂SO₄, HCl, HClO₄, H₃PO₄ and CH₃COOH) and basic media (NaHCO₃ and Na₂CO₃), resulting complex exhibits maximum and stable absorbance in phosphoric acid medium as is shown Table 1 and Figure 3.However, highly turbid solutions were obtained while using basic media for complexation. Phosphoric acid is observed as the appreciable complex forming medium and hence selected. The Ir (III)-HToC complex attained maximum absorbance (0.310) in the acidity range of 0.020-0.045M concentration of H₃PO₄. Therefore, acidity of the complexing medium is adjusted to 0.035M of H₃PO₄ (Table 2; Figure 4A).

Conditions: Ir (III) =10 μ g; Acidity of aqueous medium = 0.025M; HToC [0.1% (w/v)] in ethanol = 1ml; Solvent = water; aqueous phase = 10 ml; λ_{max} = 423 nm

Conditions: Ir (III) = $10 \mu g$; Acidity of aqueous medium (using H₃PO₄) = variable; HToC [0.1%(w/v)] in ethanol =1ml; Solvent = water; aqueous phase = 10 m l; λ_{max} = 423 n m

Effect of Reagent (HToC) concentration

The metal ion starts forming a coloured species on addition of the reagent whose absorbance in aqueous phase increases with increase in HToC content. In the range of 0.5–1.4 ml of 0.1% (w/v) ethanolic solution of reagent, the absorbance attains a maximum value of 0.310. Decrease in colour intensity results if concentration of reagent is further increased (Table 3; Figure 4B). Therefore, 1ml of 0.1% HToC is selected for further studies.

Conditions: Ir (III) = $10 \mu g$; Acidity of aqueous medium = 0.035M; HToC [0.1% (w/v)] in ethanol = variable; Solvent = water; aqueous phase = 10 m l; λ_{max} = 423 nm

Interference studies

To examine the tolerable value of the diverse ions, the foreign ion effect was studied under ideal conditions of the procedure while keeping a constant metal ion concentration (10 μg of iridium (III) in 10 mL aqueous volume) for each. The ions or complexing agents were added as their sodium or potassium salts, initially in large amounts but the reduced quantities if an error of >1% is noticed in certain cases(Table 4).Similarly, effect of cations on the colour intensity of iridium complex is also studied by taking 10 μg of iridium (III) per 10 mL aqueous phase and adding suitable amounts of different cations (Table 5). It is found that of the 33cations studied, none effected the absorbance of Ir(III)-HToC complex.

Conditions: As mentioned in procedure





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Conditions: As mentioned in procedure *Initial oxidation state shown in parentheses.

Calibration curveand statistical analysis

Iridium(III) determination employing HToC as complexing reagent depends linearly on the metal ion concentration in the aqueous solution and is narrated by the regression equation as Y= 0.315X-0.007, where X in the equation denotes concentration of Ir(III) in μg ml⁻¹(R=0.9981). it has been observed from the calibration curve that Beer's law holds good over the Ir(III) concentration range of 0.0-1.8 μg ml⁻¹(figure 5A). Beyond this concentation, it starts deviating from linearity. The optimum concentration range for accurate determination, as evaluated from a Ringbom plot [32], however, is 0.3-1.6ppm of Ir (III) (figure 5B). The standard deviation was \pm 0.0045 and relative standard deviation(RSD) = 0.0146% depicted that the developed method is precise and accurate. The method is highly sensitive as is indicated from its molar extinction coefficient as 5.960×10⁴ 1 mol⁻¹ cm⁻¹. The various analytical parameters have been placed in table 6.

Stoichiometry of Ir(III)-HToC complex

1:3 [Ir(III):HToC] ratio of metal to ligand in the developed complex is estimated by Job's continuous variations method [33,34]by mixing equimolar concentrations (1.040×10^{-3} M) of both the metal ion and reagent and measuring the absorbance at 3 divergent wavelengths 380, 423 and 450 nm as presented in figure 6A. The 1:3 complex ratio has further been confirmed by mole ratio methods [35]where in a clear break in the curves corresponding to mole ratio(reagent/metal) 3.0 suggests the metal to ligand ratio to be 1:3 (figure 6B). The above stoichiometric investigation supports the metal-ligand complex to have the optimized structure as

Computational results

Geometry optimization

After geometrical optimization, with the help of FMOs, a molecule's desired locations in the donor-acceptor connection between the ligand and the complex are precisely represented. Figure 8shows the optimized structure and electronic distribution of frontier molecular orbitals for chromogenic reagent (HToC) and its complex with Ir(III). E_{HOMO} and E_{LUMO} are generally used as reactivity markers in terms of a molecule's willingness to give and receive electrons under favourable conditions. The energy gaps for the HOMO and LUMO molecules account for the charge transfer interactions that occur within the molecule. The primary characteristic of this electronic absorption, associated with the transition from ground to the first excited state, is the single electron excitation from HOMO to LUMO. The energy gap (E_{gap}) between the HOMO (-8.28 eV) to LUMO (-5.92 eV) is 2.36 eV forthe reagent, while for complex it is determined to be 1.05 eV. HOMO and LUMO of the reagent are distributed over the whole molecule and therefore showed π - π * transition. The E_{gap} for reagent is more than that of complex.

Furthermore, the Ir(III)-HToCcomplex has a smaller dipole moment (0.359293) than that ofHToC (3.257415) indicating weak dipole-dipole interactions. Furthermore, according to other chemical descriptors such as chemical hardness/global hardness (χ), directly related to it, the HOMO-LUMO energy gap is larger for the reagent than its complex. A substance with a high chemical hardness grade is said to be stable. Similarly, the strength with which electrons are pulled to a covalent bond is determined by its electronegativity value. The inverse of global hardness, global softness serves as an additional metric to evaluate the low reactivity of the stable complex. Further, Table 7 shows that the interaction between the ligand and Ir(III) causes the chromogenic reagent's chemical hardness to decrease from 1.18 eV to 0.53 eV, indicating that the formed complex had lower kinetic stability and more chemical reactivity. Table 7 demonstrates that there was a significant charge transfer from the reagent to the metal due to its increased electro negativity compared to its metal complex. The energy stabilization that occurs when a system picks up an extra electronic charge from its surroundings, is measured by the electrophilicity index. The complex and ligand had electrophile and nucleophile properties, respectively, according to the computed electrophilicity index. Ultimately, the desire of a molecule to provide and receive electronic charge is known as its capacity for giving and receiving electrons, respectively.





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QTAIM Study

By using Bader's topological QTAIM analysis, we examine the type and intensity of metal-ligand interactions [36]. The reagent(HToC) and Ir(III)-HToC complex's molecular graph is graphically represented in figure 9, showing the locations of bond critical points (BCPs) and the bond pathways that connect the oxygen atoms (O11, O12, O20, O21, O27, and O34) of the ligand with iridium metal. Table 8 provides topological metrics at BCPs of intermolecular interactions between the reagent and metal, including electron density (QBCP), laplacian (∇^2 QBCP), total energy density (HBCP), kinetic energy density (GBCP), and potential energy density (VBCP). Additionally, the sign of the HBCP and ∇^2 QBCP values can be used to determine the intensity of the intermolecular interactions. Thus, a strong interaction (∇^2 QBCP and HBCP<0), a medium strength interaction (∇^2 QBCP and HBCP<0), and a weak interaction (∇^2 QBCP and HBCP>0) may be distinguished. The findings presented in Table 8 indicate that the metal-ligand interaction strength is medium, as indicated by the positive values of ∇^2 QBCP and the negative values of HBCP for Ir-O bonds. A good criterion for figuring out the type of interaction is the ratio of GBCP/|VBCP| [37]. Covalent, electrostatic with partially covalent, and non-covalent interactions are shown by the ratios of 0.5 < GBCP/|VBCP| > 0, 1 < GBCP/|VBCP| 1, and so on. The computed GBCP/VBCP value for Ir-O bonds is found to range between 0.5 and 1, as indicated in the Table 8. This indicates that the interactions are primarily electrostatic with a partial covalent nature.

Applications

Analytical Applications

For the micro amount of Ir (III), the proposed technique of spectrophotometric determination is simple, rapid and offers the advantage of high sensitivity ($\varepsilon = 5.960 \times 10^4 \, l \, mol^{-1} cm^{-1}$) and selectivity (free from interference of 23 anions/complexing agents and 33 cations of analytical interest). The wide practicability of the procedure has been assessed by analysing different samples of varying compositions and the results obtained are quite satisfactory(Table9). The total operation for each determination requires not more than 5minutes.

- *Figure in parantheses indicates the amount of metal ion in mg
- ** Average of duplicate analyses

Bioanalytical applications

DPPH radical scavenging capacity

The management of antioxidants play a vital role for the preservation of biological systems. Scavenging free radicals and inhibiting chain reactions work as antioxidants by providing the free radical. Safer antioxidant has been created in the presented work. The antioxidant capacity at various concentrations of HToC and Ir (III) complex was analysed by detecting the de-colorization of DPPH. %RSA increases with enhanced concentrations of the compound under study. HToC and Ir (III)-HToC complex exhibited the highest scavenging activity of 59.3% and 70.04%, respectively, at 500 μg ml $^{-1}$ concentration. It has also been noticed that the Ir (III)-HToC complex, a chelate, acted as better antioxidant as compared to reagent alone. The RSA can be ranked in the following order: gallic acid > Ir (III)-HToC complex > HToC. This indicates that the prepared complex has the potential to be highly influencial in the development of innovative, therapeutic approaches that rely on antioxidants.

CONCLUSION

In the reported work, spectrophotometric determination technique has been used to analyse iridium(III) at the microgram level using a new derivative from the family of benzopyrans, 3-hydroxy-2-tolyl-4H-chromen-4-one(HToC) aschromogenic reagent. In phosphoric acid medium, iridium (III) instantly interacts with HToC to generate a stable light yellow-coloured, non-extractable complex that coherently responded linearly upto 1.8 Ir (III) µg ml⁻¹. The corresponding values of the statistical parameters viz. %RSD, molar absorptivity, Sandell's sensitivity and corelation coefficient, validated the method's precision and accuracy. The procedures under study confirmed that the produced complex had stoichiometric constitution of 1:3[Ir (III):HToC].The molecular structural features are elucidated by TDDFT analysis and QTAIMevaluation. The method found its satisfactory importance in





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numerous analytical applications in different synthetic mixtures of varying compositions. The antioxidant findings of the complex reveal its best potential to scavenge radicals.

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Table 1: Effect of medium on the absorbance of Ir (III)-HToC complex

Medium U	sed	Absorbance	Medium Used	Absorbance
H ₃ PO ₄	H ₃ PO ₄ 0.310		HCl	0.118





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HNO ₃	0.119	H ₂ SO ₄	0.176
CH ₃ COOH	0.085	HClO ₄	0.126

Table 2: Effect of acidity on the absorbance of Ir (III) –HToC complex

H ₃ PO ₄ (M)	0.005	0.010	0.015	0.020-0.045	0.05	0.06	0.075
Absorbance	0.09	0.19	0.293	0.310	0.255	0.210	0.175

Table 3: Effect of concentration of reagent on the absorbance of Ir (III)-HToC complex

Amount of HToC added (ml)	0.1	0.3	0.4	0.5-1.4	1.5	1.7	2.0
Absorbance	0.020	0.160	0.200	0.310	0.280	0.127	0.080

Table 4: Effect of anions or complexing agents on Ir(III)-HToC complex

Anions/ Complexing agent	Tolerance limit	Anions/ Complexing agent	Tolerance limit
added	mg/10mL	added	mg/10mL
Chloride	100	Phosphate	100
Bromide	100	Thiourea	100
Iodide	100	Sulphate	90
Nitrate	100	Ascorbic acid	90
Nitrite	100	Acetate	50
Sulphite	100	Fluoride	50
Carbonate	100	Oxalate	50
Bicarbonate	100	Dithionite	50
Sulfosalicylic acid	100	Tartrate	40
EDTA'Disodium salt'	100	H ₂ O ₂ (30%)	1ml
Hydrazine sulfate	100	Glycerol	1ml
Thiocynate	100		

Table 5: Effect of cations on Ir (III)-HToC complex

Table 3. Effect of Cations on it (III)-1110C complex								
Cations added*	Tolerance limit mg/10mL	Cations added*	Tolerance limit mg/10mL					
Cu(II)	10	Pd(II)	1					
Zn(II)	10	Fe(II)	1					
Mg(II)	10	Fe(III)	1					
Mn(II)	10	Os(VIII)	1					
Ca(II)	10	Zr(IV)	1					
Sr(II)	10	Sn(II)	0.8					
Ni(II)	10	W(VI)	0.5					
Co(II)	10	Ti(IV)	0.5					
Cr(III)	10	Au(III)	0.5					
Cd(II)	10	V(V)	0.5					
Se(IV)	8	Mo(VI)	0.5					
Al(III)	7	Ru(III)	0.5					
Ba(II)	7	Pt(IV)	0.5					
Ce(IV)	5	Nb(V)	0.5					





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Pb(II)	5	As(V)	0.5
Ag(I)	5	Cr(VI)	0.1
Hg(II)	1		

Table 6: Analytical characteristics of Ir (III)-HToC complex

S.NO.	Characterstics	Specification
1	λ_{max}	420-425 nm
2	Molar absorptivity(ε)	5.960×10 ⁴ l mol ⁻¹ cm ⁻¹
3	Sandell's sensitivity(S)	0.0032 μg cm ⁻²
4	Beer's law range (B.L.)	0.0-1.8 μg ml ⁻¹
5	Ringbom's ideal range	0.3-1.6 μg Ir (III) ml ⁻¹
6	Linear regression equation	Y = 0.315X - 0.007
7	Correlationcoefficient	0.9981
8	Limit of detection (LOD)	0.036 µg ml⁻¹
9	Standard deviation	0.0045
10	%Relative Standard deviation	0.0146
11	Coefficient of variance	0.00146
12	Stoichiometry of the complex	1:3 (M:L)
13	Stability of the complex	>2 days

Table 7: Calculated HOMO-LUMO energy gap (Egap) and quantum molecular descriptors in eV for the studied reagent and complex Ir(III)-HToC

studied reagent and complex in (iii) irroc									
Parameters studied	Еномо (eV)	Elumo (eV)	Egap (eV)	I	A	χ (eV)	η (eV)	σ	ω
НТоС	-8.28	-5.92	2.36	8.28	5.92	7.10	1.18	0.85	21.37
Ir(III)- HToC complex	-5.89	-4.83	1.05	5.89	4.83	5.36	0.53	1.90	27.29

I= Ionization potential; A= Electron affinity; χ = Electronegativity; η = chemical hardness; σ = chemical softness; ω = electrophilicity index HToC

Table 8: Evaluated topological parameters (a.u) at BCP in the studied Ir(III)-HToC complex

Bond	Q BCP	$\nabla^2 Q$ BCP	Нвср	GBCP	$\mathbf{V}_{\mathtt{BCP}}$	$G_{BCP}/ V_{BCP} $
Ir-O12	0.09334	0.5323	-0.0085	0.1380	-0.1466	0.94133697
Ir-O11	0.0937	0.5047	-0.0099	0.1326	-0.1425	0.93052632
Ir-O34	0.0975	0.5296	-0.0105	0.1391	-0.1496	0.92981283
Ir-O27	0.0936	0.5337	-0.0086	0.1384	-0.1470	0.9414966
Ir-O21	0.0939	0.5077	-0.0099	0.1333	-0.1432	0.93086592
Ir-O20	0.0894	0.5031	-0.0081	0.1306	-0.1387	0.94160058

Table 9: Analysis of synthetic mixtures by the studied method

	Composition of sample		Ir found**	
S.No.	Matrix *	Ir added (μg / 10 ml)	(μg / 10 ml)	
1	Cu(5), Zn(2), Se(0.1)	10	10.05	
2	Mg(2), Mn(5), W(0.5)	5	4.95	
3	Ca(5), Ni(2), Pb(0.1)	15	15.11	





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4	Co(5), Cr ^{III} (2), Mo(0.1)	12	12.06
5	Zn(5), Pt(0.5), V(0.1)	7	7.08
6	Os(0.1), Ce (1), Fe(0.5)	10	10.07
7	Zr(0.5), Hg(0.1), Ti(5)	8	7.90
8	Ca(5), Cd(0.5), Ba(2)	5	4.95
9	Mn(5), Fe(0.1), Ag(0.1)	7	7.50
10	Mg(2), Sn(0.1), Pd(0.1)	12	11.94

Table 10: %RSA of Ir(III)-HToC complex

Concentration	%RSA			
(µg ml⁻¹)	HToC#	Ir(III)-HToC##	Gallic acid###	
31.2	34.8	37.2	47.6	
62.5	38.1	42.5	52.2	
125	45.8	58.1	65.7	
250	54.5	65.1	74.3	
500	59.3	70.4	80	

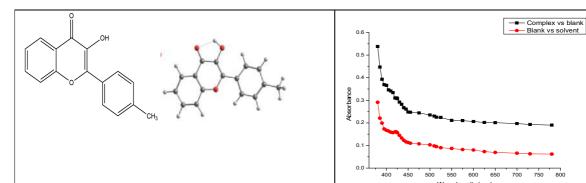


Figure 1. Structure of HToC

Figure 2. Absorbance spectrum of Ir (III)-HToC complex $\,$

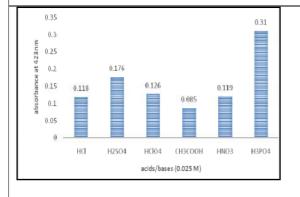


Figure 3. Effect of medium on the absorbance of Ir (III)–HToC complex

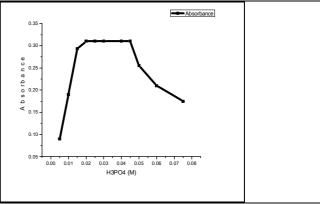


Figure 4. Effect of acidity[A] and reagent concentration

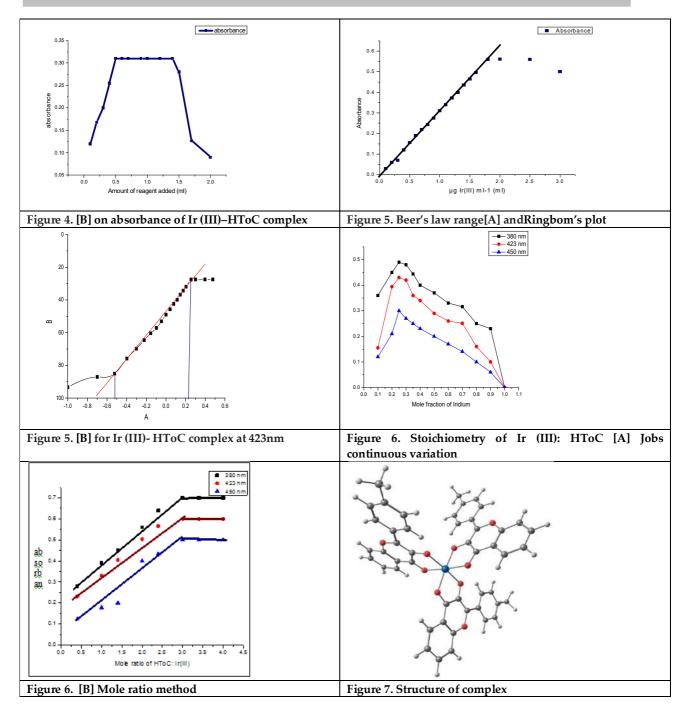




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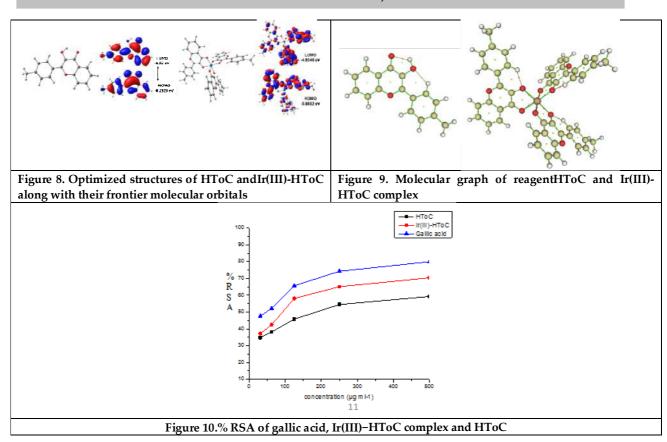
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REVIEW ARTICLE

Review of Covalent Organic Frameworks and their Applications

Shipra Sharma, Satvinder Khatkar, and Bhawna Pareek*

Department of Chemistry, Maharishi Markandeshwar (Deemed to be University), Mullana, Ambala, 133207, Haryana, India.

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*Address for Correspondence

Bhawna Pareek

Department of Chemistry,

Maharishi Markandeshwar (Deemed to be University),

Mullana, Ambala, 133207, Haryana, India.

Email: dr.bhawnapareek@gmail.com,



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ABSTRACT

Covalent Organic Frameworks are two or three dimensional (2D or 3D) structures in which monomers are connected by covalent bonds. The extended crystalline solids can be formed by linkage of organic units by covalent bonds between light elements like (B, C, N, H, O) to form covalent organic frameworks (COFs). Gilbert N Lewis in 1916 describe the covalent bond and nature of bonding within the molecules and linkages of atom to form molecules. In 2005 Omar M. Yaghi and his coworkers discovered the first crystalline covalent organic framework COF-1. Different methods are used to make crystalline covalent organic frameworks like room temperature synthesis, solvothermal methods, microwave synthesis, ionothermal methods of synthesis etc. The most common method of synthesis is solvothermal method which is used for the formation of highly crystalline and stable covalent organic frameworks. Covalent organic frameworks are highly ordered crystalline porous structures having combination of both physical and chemical properties. The band gap of COFs can be tuned by different molecular building blocks. Due to their strong covalent bonds, high porosity, high solubility, high crystallinity and tunability COFs can be used in Hydrogen storage, Methane storage, Gas separation, Sensing, Catalysis, drug delivery system, CO2 conversion etc. Covalent organic frameworks and COFs linked hybrid materials are very important in photodegradation of dyes and pollutants.

Keywords: Covalent Organic Frameworks, Covalent Bonds, Photodegradation, Crystalline porous structures etc.





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INTRODUCTION

Covalent organic frameworks are highly crystalline material having two dimensional (2D) or three dimensional (3D) porous crystalline structures having high surface area and high thermal stability. Covalent organic framework has low density which is due to less dense elements present in organic linkers connected by strong covalent bonds. The first COF material was discovered by Yaghi and his coworkers in 2005 i.e COF-1. Covalent organic framework an emerging crystalline material having high porosity and crystallinity with tunable pore size and having large self-healing capacity. Covalent organic frameworks (COFs) are playing an important role in research now days due to their combined physical and chemical properties. Different methods are used in synthesis of COFs and number of methods have been developed for improving band structure and photocatalytic activities of Covalent Organic Frameworks.

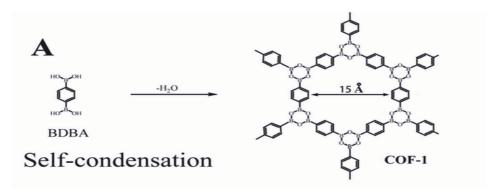
Linkages

The building blocks of Covalent Organic Frameworks are made up of different linkages having Boron, Nitrogen, Sulphur and Oxygen having strong covalent bonds present in their molecular building units. These linkages introduced electron donor acceptor unit into conjugated system to extend light absorption towards visible light. The different building blocks of linkages are given below. (6)

- 1. Boron containing linkages
- 2. Triazine based linkages
- 3. Imine based linkages
- 4. Imide based linkages
- 5. Hydrazone based linkages
- 6. Azine based linkages

Boron containing linkages

Boroxine linkage, Boronate ester linkage, Boronated anhydride linkage can be synthesized by self condensation of Boronic acid(RBOH2) or co-condensation of boronic acid (RBOH2). Boron containing COFs are highly sentient of water and moisture so can be easily oxidized and hydrolyzed. Boron containing linkages have low density but have high photoconductivity, high crystallinity and large surface area and high thermal stability as compared to others linkages. Examples are COF-1.



Triazine based linkages

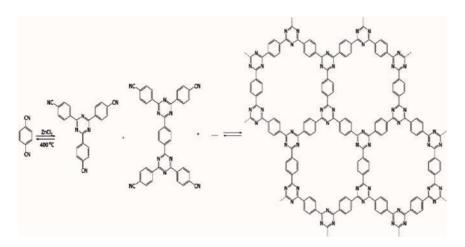
Covalent triazine based frameworks (CTFs)which can be synthesized by cyclo-trimerization of nitriles under ionothermal condition and Friedel craft reaction. Triazine based COFs have superior stability as compared to boron-containing linkages but have low crystallinity due to presence of large





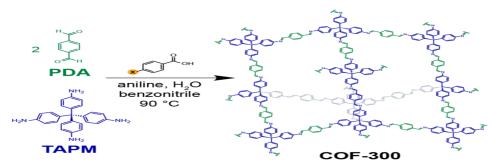
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number of Nitrogen atoms and high degree of conjugation and high structural tunability e.g. CTF-1



Imine based linkages

Imine based COFs are synthesized by co-condensation of amine with aldehydes in the presence of Lewis acid catalyst or an organic acid. Imine linked COFs show more stability in water, acidic or basic condition and organic solvent but inferior crystallinity. Imine liked COFs show low pi-delocalization due to high polarization of Nitrogen atom. Imine linked COFs can chelate with metal ion due to presence of large no of nitrogen atom e.g. 3D COF-300



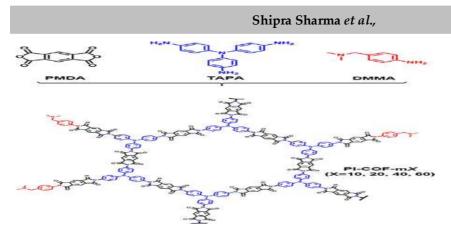
 $X = OH, OCH_3, CH_3, H, F, CI, Br, I, CF_3, CN, NO_2$

Imide based linkages

Imide linkage can be synthesized by condensation of amine with acetic anhydride up to 250° C temperature A series of crystalline polyimide COFs were fabricated via reversible imidization reaction by extending the building blocks and the large pore size of PI-COFs could be tuned. e.g. PI-COFs-1

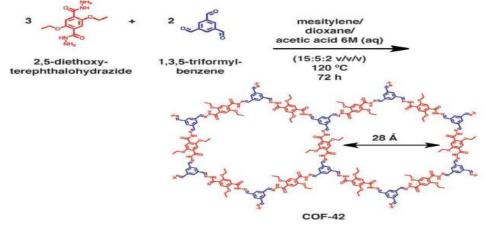






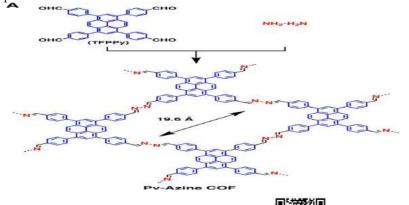
Hydrazone based linkages

Hydrazone linked COFs have been synthesized by co-condensation of aldehydes and hydrazides in presence of acetic acid as catalyst. Due to weak interlayer interactions in hydrazone linked COFs they can be exfoliated into thin film under mild conditions E.g. COF-42.



Azine based linkages

Azine based COF have linkage present in most of COF. It is result of short hydrazine monomer being used to connect two aldehydes form polygon skeleton. The azine linked COFs have permanent porosity and high surface area and high chemical stability. E.g. Py-Azine COF. Azine based covalent organic frameworks are used as metal free visible light photocatalyst in carbon di oxide reduction and hydrogen production.





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Methods of synthesis

Different methods for the synthesis iof Covalent Organic frameworks powders are Solvothermal Method, Ionothermal, Room temperature, Microwave synthesis.(6)

Solvothermal Synthesis

Most of the COFs are synthesized by Solvothermal Method. In this Method reaction is carried out under heating (80 to 120 °C) in sealed vessel for 3 to 7 days. The selection of solvent is very important like 1,4 –dioxane/mesitylene), DMAC-o-Chlorobenzene have been utilized to fabricate COFs liked with organic linkers to fabricate crystalline framework in COF. Solvothermal method of synthesis is very time consuming process as the reaction rate of COF synthesis is very slow.

Ionothermal Method

In Ionothermal method Molten metal salt or ionic liquids are used as both solvent and catalyst to form solid at high temperature around 400°C and high pressure. Harsh condition like high temperature(around 400°C) and high pressure limits the application of Ionothermal method. e.g COFs prepared by Ionothermal method are Covalent Triazine framework.

Room Temperature Synthesis

Room temperature synthesis consist of two methods

- a) Mechanochemical Grinding
- b) Solvent Method

Mechanochemical Grinding

This method is used by pestle and mortar to prefer manual grinding. It is simple, rapid, facile, environment friendly, solvent free method carried out at room temperature. The change of colour indicate the formation of COFs.

Solvent Method

In this method liquid assisted mechanical grinding was used in which addition of catalyst solution increases the rate of chemical reaction.

Microwave assisted Synthesis

Microwave assisted method of synthesis has short reaction time, high product yield, low energy usage, cleaner products and controllable reaction temperature and pressure and 200 times faster than solvothermal method of synthesis and has large BET surface area as compared to solvothermal method.

Applications

COFs can be used in Hydrogen storage, Methane storage, Gas separation, Sensing, Catalysis, drug delivery system, CO₂ conversion etc.(2)

Drug delivery

Drug delivery using Covalent organic frameworks spotlight the encapsulation and decapsulation of drugs in a versatile way. Covalent organic frameworks are the good drug carriers and delivers drug to the target molecules in effective way which solve the many drug delivery problems.





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Sensing

Covalent organic frameworks are used as a electrochemical sensor as well as biological sensors due to their unique structures and except physical and chemical properties. Detecting chemicals and biomolecules in the fields of environmental protection, health monitoring and for public benefits a large numbers of chemical and biomolecules are detected by detectable signals which can be electrical, thermal by the selective adsorption of analytes.

CO₂ Conversion

Covalent organic frameworks have essential role in carbon di oxide reduction and conversion. COFs could convert the CO₂ in high valuable product which is not harmful for environment. As COFs can transform the CO₂ into HCOOH, CH₃OH at different redox potentials.

Hydrogen production

Covalent organic frameworks can be used as the production and evolution of hydrogen through water splitting by using method photocatalysis and electrocatalysis which are highly dependent of different structure of COFs.

Gas Storage

Covalent organic frameworks have large numbers of pore structures and large surface area which further increses the storage of gas molecules in it. Also the tunable pore size increses their importance in the gas storage process as well as gas separation process.

Catalysis

COFs based photocatalytic reaction show superior photocatalytic activities which show their possible application in development of environmental friendly reaction using visible lights well as ultraviolet light. Photocatalysis of COFs shows many photocatalytic degradations of dyes, pesticides, and environmental pollution.

CONCLUSION

COFs are the class of highly porous crystalline materials having high photocatalytic properties and structural versatility and tunability which shows their large capacity to degrade aqueous pollutants like dyes, pesticides, antibiotics under visible light or sun light irradiation. COFs are useful in degradation of various pollutants but adsorption of pollutants is still under exploration. But literature review presented various photocatalytic application of COFs in water splitting, CO2 reduction, gas storage, degradation of aqueous pollutants and drug delivery system but still there are many challenges in formation of crystalline covalent organic frameworks and also to maintain the stability of COFs.

Declaration of Interest

The author declared that they do not have any conflict of interest.

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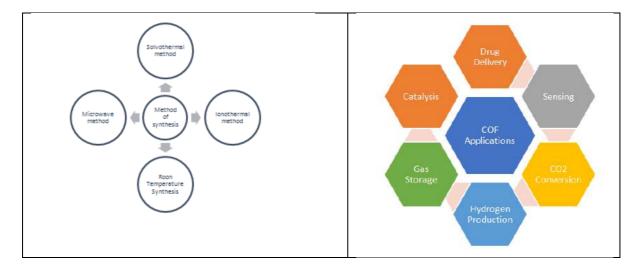
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RESEARCH ARTICLE

Catalytic role of Carbon Quantum Dots in Super Capacitor Applications Synthesis, Characterization, and Properties

Karamveer Sheoran and Samarjeet Singh Siwal*

Department of Chemistry, M.M. Engineering College, Maharishi Markandeshwar (Deemed to be University), Mullana-Ambala, Haryana, 133207, India.

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*Address for Correspondence Samarjeet Singh Siwal

Department of Chemistry,

M.M. Engineering College,

Maharishi Markandeshwar (Deemed to be University),

Mullana-Ambala, Haryana, 133207, India.

Email: samarjeet6j1@gmail.com



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ABSTRACT

Carbon quantum dots (CQDs) have emerged as promising materials in various technological applications due to their unique physicochemical properties. In the realm of energy storage, their role in super capacitors has garnered significant attention. This paper investigates the catalytic role of carbon quantum dots in enhancing the performance of SCs. The synthesis methodologies of CQDs and their subsequent characterization techniques are elucidated, emphasizing their structural, morphological, and surface properties. The electrochemical behavior and specific capacitance of SCs employing CQDs as active materials are thoroughly examined. The superior charge storage capability, high conductivity, and remarkable stability exhibited by CQD-based SCs are discussed in detail. Additionally, insights into the mechanisms underlying the enhanced electrochemical performance facilitated by CQDs are provided. This comprehensive analysis underscores the potential of CQDs in revolutionizing the development of efficient and high-performance super capacitors for advanced energy storage applications.

Keywords: Carbon quantum dots; Energy storage applications; Electrochemical performance; Physicochemical properties.

INTRODUCTION

The demand for sustainable energy supply, it's storage and environmental significance have been surging worldwide. Owing to unique features such as high power density, excellent cycle stability, elongated life cycle, and cost-effective, super capacitors (SCs) emerged as potential energy storage devices which grabs the focus of scientific





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communities [1, 2]. However, the relatively low energy density restricts the widespread application of SCs. Multitude efforts have been made by researchers towards the development of high energy density SCs but it still remains challenging. To boost the energy density of SCs, researchers can inflate either the specific capacitance or the voltage window. Nevertheless, the electrocatalytic performance of SCs are greatly depends on the nature and the type of materials for their electrodes and electrolytes[3]. Therefore, the electrochemical activity of the SCs can be improved with the fabrication of novel electrode materials. With regard to this, carbon quantum dots (CQDs) have arisen as a novel class of nanomaterials that hold great promise for SC applications. CQDs possess exclusive characteristics such as excellent biocompatibility, high stability, and ultra small particle size (<10 nm) resulting not only from quantum confinement but also due to edge effects and surface passivation [4, 5] which makes them suitable candidate for energy storage devices. CQDs can be utilized as electrolyte additives, active materials and conductive additives in SCs to enhance electrolyte features, electrical conductivity and to store energy respectively. The integration of CQDs into SC materials offers opportunities for improving the electrochemical performance and energy storage capabilities of SCs. This review summarized and discussed the potential of CQDs in revolutionizing the development of efficient and high-performance supercapacitors for advanced energy storage applications. We systematically elucidate synthetic approaches of CQDs along with different properties and catalytic role of CQDs in SCs. Recent research findings and case studies on CQDs-based SCs are presented to provide insights into their performance and potential applications. Finally, the challenges and future prospects of CQDs and it's composites for SC applications are discussed.

Synthesis methods of carbon quantum dots

Usually, CQDs have been synthesized with the utilization of "top-down" and "bottom-up" approaches where high dimensional carbon materials like graphene, graphite, carbon nanotubes, and graphene oxides (GO) are used as starting materials in case of top-down approach for the synthesis of CQDs by physical or chemical techniques whereas smaller carbon materials are utilized in case of bottom-up approach for the fabrication of CQDs by chemical reactions. Different top-down and bottom-up approaches are demonstrated in **figure 1**.

Top-down approach

Laser ablation, arc-discharge, ultrasonic, chemical exfoliation and electrochemical methods have been utilized in top-down approach for the synthesis of CQDs. In this section, we briefly discussed these techniques to fabricate CQDs.

Laser ablation

Laser ablation is widely used method for the fabrication of CQDs with diverse morphologies. This method was firstly purposed by Sun *et al.*[6]with the utilization of Nd:YAG laser source (1064 nm, 10 Hz) for the irradiation of carbon raw material followed by acid treatment. The obtained CQDs did not emit light initially; but when the surface was modified with two different oligomers i.e., poly(ethylene glycol) and poly(propionylethylenimine-coethylenimine), CQDs exhibited strong photoluminescence emission. After that CQDs of different sizes have been synthesized successfully under controlled nucleation. At the interface of solid target and surrounding liquid media of target, plasma pulse can be localized by using a high energy laser. This plasma pulse initiates rapidly expandable bubble. This bubble shrinks after the application of pressure from surrounding liquid and the nucleation of CQDs was slowly started by cooling core. With the adjustments of laser pulse width, CQDs with different cluster densities can be developed[7].CQDs were prepared by Li *et al.* [8] using simple laser ablation technique where dispersion of nano-carbon material in 50 mL of solvent was used as starting material. Nd:YAG pulsed laser was used for laser irradiation to obtain CQD. CQDs with narrow particle size distribution and outstanding fluorescence features can be fabricated by using laser ablation method. Though high cost and complicated process limits it's application in several fields.

Arc-discharge

In the arc-discharge method bulk carbon precursor undergo decomposition in anodic electrode in the presence of plasma generated in a sealed reactor[9]. The temperature within the reactor reaches up to 3727 °C, where vaporisation





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of the carbon atoms occurred at anode and then reassembled at cathode which results in the formation of CQDs. Without the modification of surface, the obtained CQDs have higher oxygen content and highly fluorescent characteristics[10]. However, due to the presence of complex impurities, purification of CQDs produced by this technique is difficult [9].Xu *et al.*[11]prepared CQDs accidently while synthesizing single-walled carbon nanotubes (SWCNTs) with the utilization of arc discharge technique. At 365 nm, yellow, blue-green, and orange fluorescence were emitted by the as-prepared CQDs. The reports showed that the CQDs fabricated with the utilization of this method exhibits good water stability but have larger particle size distribution because of which the surface area of CQDs was decreased extensively and restrict electrocatalytic performance of the material.

Ultrasonic

Ultrasonic techniques are considered as convenient and simple method for the fabrication of CQDs. In this approach, low-and high-pressure waves are created in the solution by high-intensity ultrasound that generate collapsing bubbles with high temperatures(5500 K) and pressure (500 atm.). For the production of CQDs, high energy spot act as micro reactors for dehydration, polymerization and carbonization and then single nuclear burst of precursors have been used[12]. A team of researchers synthesized Sn@CQDs@Sn via ultrasound method. The authors observed that the as-prepared CQDs are stable up to 90 cycles and reached maximum coulombic efficiency of 99% for energy storage device [13].

Chemical exfoliation

The electrochemical cleavage of different carbon materials like graphene, reduced graphene oxide, graphite rods, and CNTs results in the production of CQDs and this approach to synthesize CQDs is known as chemical exfoliation technique. In this technique, OH and O are produced at the anode by the oxidation of water. These radicals work as electrochemical scissors to generate CQDs. The process of exfoliation generally occurs at edge points and become faster at the defective points[14]. This method is considered as promising facile and approach for the large scale formation of CQDs as no harsh conditions and strong acids are needed in this technique [15].

Electrochemical Methods

Electrochemical method is prominent and convenient method to fabricate CQDs owing to several merits such as abundance of raw materials, cost-effectiveness, and facile preparation under normal conditions of temperature and pressure. CQDs were prepared by Chi *et al.*[16] using electrochemical method using phosphate buffer solution (pH 7.0) where working electrode was graphite electrode, and Pt mesh and Ag/AgCl was counter and reference electrode respectively. The only demerit of this technique is it's tedious purification process.

Bottom-up approach

Hydrothermal, microwave and thermal pyrolysis techniques are involved in bottom-up approach for the fabrication of CQDs. Saccharides and citric acids are generally utilized as building blocks to prepare CQDs. With the modifications of experimental conditions, CQDs of desired size can be obtained in cost-effective way. This section briefly elucidates bottom-up approaches to fabricate CQDs.

Hydrothermal method

Hydrothermal technique of CQDs preparation involves the carbonization of organic precursors which are sealed under high temperature of up to 200 °C in Teflon-lined hydrothermal reactor. Carboxylic acid and amino groups were utilized as precursor for the formation of CQDs. Further studies on hydrothermal synthesis of CQDs revealed that monopotassium phosphate and glucose can also be utilized as precursor to prepare CQDs. For instance, Vandana and co-workers [17]synthesized CQDs via hydrothermal method by utilizing glucose as precursor. The assynthesized CQDs were used as electrode material for SC and exhibited excellent specific capacitance of 647.5 F/g with decent energy density and power density of 93 Wh/kg and1430 W/kg respectively. Furthermore, the assynthesized CQDs possess high stability of 2000 cycles.





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Microwave method

In microwave synthesis, electromagnetic radiations of 1mm to 1 m wavelengths allow the interaction with precursor and results in CQDs generation within 10 min[18]. The CQDs formed via microwave method exhibits several surface functional and broad size distributions because of abrupt heating of microwave resonance. CQDs were prepared by Kumar *et al.*[19]synthesized CQDs via microwave technique using solution of citric acid and urea as precursor. The CQDs were prepared within 300 sec. This technique is facile and scalable method but poor control over obtained particle size restricts the performance of CQDs in various fields.

Comparative analysis of synthesis methods and their impact on CQD properties

The synthetic methods for the formation of CQDs have some pros and cons which are summarized in table 1.

Properties and catalytic role of CQDs in supercapacitors

The electron transfer features of CQDs are regulated by the interactions between carbogenic core and functional groups. High surface area and available active sites of CQDs enhances the electron transfer process. The enhancement in the catalytic features of CQDs can be obtained with the doping of heteroatoms like S, B and N etc. The oxygenated functional groups present on edges also enhances the catalytic feature of CQDs[21]. This section summarizes the beneficial property and utilization of CQDs in SCs.

Electrochemical properties of carbon quantum dots

Electron transfer properties of CQDs can be regulated by the interactions of functional groups and carbogenic core. Doping of heteroatoms also influences electron transfer of CQDs [22]. The higher surface area and availability of profuse edge sites of CQDs facilitates the transfer of electrons. Particularly, abundant functional groups which are present on CQD's surface solidly nucleate and modify the pristine nanocrystals which results in intense electrostatic interactions. Consequently, stability and electrochemical activity of the material enhances due to the strengthening of contact between active material and conductor. CQDs can be combined with other materials to enhance the electrochemical performance of the material. For example, Zheng et al. [23] prepared molybdenum disulphide (MoS2) based electrode materials and observe the electrochemical performance of modified electrodes. Briefly, the MoS2 electrode was firstly modified with zinc sulphide (ZnS); (MoS₂@ ZnS) and then modified with CQDs by utilization of facile hydrothermal method. The obtained results indicate that CQDs/ MoS2@ ZnS showed better electrocatalytic activity with specific capacitance of 2899.5 F/g and capacitance retention of 76.1% over 3500 cycles while the obtained specific capacitance of MoS₂, ZnS and MoS₂@ ZnS was 1067.6 F/g, 423.4 F/g and 2176.1 F/g respectively. In another study, CQDs/NiAl-layered double oxide (CQDs/NiAl-LDH) composite nanosheet were prepared via one step solvo thermal method [24]. The as synthesized composite exhibited pore size and surface area of 7.1 nm and 77.9 m²/g respectively. The authors revealed that the introduction of CQDs enhances the electrochemical performance of the composite by comparing the specific capacitance of their material with previously reported NiAl-LDH based materials. The as-prepared material hadspecific capacitance of 1794 F/g at current density of 2 A/g. It also showed high cycling stability of 1500 cycles with retention of nearly 93%. These results further proves that the special features such as porous structure, active sites, high charge transfer capacity, and easy accessibility to electrolytes of CQDs helps to achieve high stability, conductivity or electrochemical performance of the material which are highly beneficial in the field of supercapacitors.

CQDs in electrocatalysis

High surface area and fast electron transfer feature of CQDs make them suitable candidate for energy conversion and storage devices. To design multi component electrocatalysts using CQDs is easy because of the ease of surface modification or functionalisation. The modification of surface with different functional groups promotes the electron transfer process which in turn improves the electrocatalytic activity of CQDs[21]. Samantara and team [25] prepared CQD composite (rGO/N-S-CQD) which was used as electrode material for SC and observed that the as-prepared CQDs exhibited decent energy and power density having cyclic stability over 5000 cycles. This showed that the doping of hetero atoms significantly increases the number of active sites for reactions of electrocatalytic devices.





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CONCLUSION, EXISTENCE CHALLENGES AND FUTURE PROSPECTS

CQDs are considered as potential candidates for the application of next generation SCs. This review provides different synthetic approaches of CQDs, their electrocatalytic properties and the role of CQDs in electrocatalysis. Despite of numerous advantages of CQDs in SCs, the research on CQDs in SCs is still in infancy stage which indicates that several avenues are yet to explored. There are still plentiful chances to use CQDs and their composites in SCs. There are some specific challenges that need to clarified in near future for the advanced research of CQD based SCs. The challenges include the deep understanding of charge storage mechanism of CQDs and diversification of functionalization and purification of CQDs. By resolving these issues industrialization of CQDs in SCs can be achieved.

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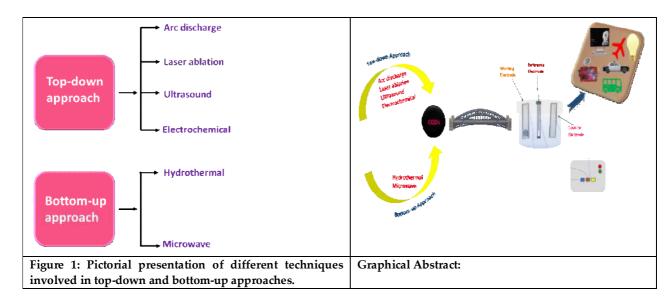


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Table 1:

Top- down Approach	Technique	Advantages	Disadvantages	
	Arc- discharge	CQDs with high oxygen content, high fluorescent	Costly, difficulty in purification and high variation in surface area	[10]
	Laser ablation	Controlled morphology, high water solubility	Process complications and high cost	[20]







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RESEARCH ARTICLE

Spatial analysis of pH, EC and TDS levels in supply water through ArcGIS mapping in different blocks of Ambala district, in North India

Saloni Kamboj¹, Nirankar Singh^{1*} and Brajesh Saxena²

¹Department of Chemistry, Maharishi Markandeshwar Engineering College, Maharishi Markandeshwar (Deemed to be University), Mullana, Ambala-133207, India.

²ICAR-Central Soil Salinity Research Institute, Karnal-132001, India.

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*Address for Correspondence

Nirankar Singh

Department of Chemistry,

Maharishi Markandeshwar Engineering College,

Maharishi Markandeshwar (Deemed to be University),

Mullana, Ambala-133207, India.

Email: nirankar singh11@yahoo.com



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ABSTRACT

Groundwater quality is a major concern in various regions around the globe, and research on quality parameters is particularly important for irrigation and drinking purposes. A systematic study of selected water quality parameters was conducted in several blocks of the Ambala district in the winters and summers of 2021–2022. Arc GIS mapping software was utilized for the spatial analysis of water quality parameters including pH, EC and TDS. Globally, water resources are investigated in conjunction with the growth of agricultural practices, industrialization, urbanization, etc. The primary goal of this study was to employ Geographic Information System (GIS) software to map the supply water quality in the studied region. In order to create spatial distribution maps of water quality parameters including pH, EC, and TDS for Ambala district, twenty predefined locations were selected from the different blocks of Ambala district, for the collection of supply water samples. These samples were then subjected to quantitative analysis and results obtained were utilized for the ArcGIS mapping.

Keywords: GIS, Ground Water, Chemical Parameters, Mapping.

INTRODUCTION

Regardless of caste, religion, nationality, gender, or social standing, everyone has the human right to access clean drinking water (Mukherje et al., 2020; Satpathy et al., 2022). However, groundwater resources are depleting globally due to rapidly growing population urbanization and industrialization(Sajjad et al., 2023; Zahran et al., 2023). Tainted drinking water can have an adverse effect on health and lead to several illnesses like diarrhea, cholera, and typhoid





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(Yasmeen et al., 2023; Misra et al., 2023). Comprehension of the current state of drinking water conditions is a fundamental prerequisite for any urban or rural area to get ready for healthy ecosystem sustainable growth(Shutaleva et al., 2020; Dutta et al., 2022). The appropriateness of ground water for drinking and other domestic uses has been investigated by various researchers using a variety of approaches and procedures (Gaagai et al., 2022). Contamination of groundwater and the variety of pollutants affecting water supplies are among the most significant environmental challenges of present time (Singh et al., 2022; Ismanto et al., 2023).

Although the inherent chemical quality of groundwater is generally found good, but various anthropogenic factors contribute to the high quantities of various elements (Saqib et al., 2023). The quality of groundwater can be significantly altered by heavily irrigated agricultural flows into the groundwater (Bodrud-Doza, et al., 2020; Balogun et al., 2022; Singh et al., 2023) An evaluation of the supply water quality has been attempted in this paper with reference to current state of pH, EC and TDS levels in the area. To ensure the quality of supply water for domestic, agriculture and industrial purposes, several parameters including the levels of pH, EC and TDS, are evaluated, and the results are compared to the acceptable/desirable values specified by the World Health Organization (WHO) and the BIS (Mazinder Baruah., 2022; Singh et al., 2024; Khapraet al., 2023). The present study offers preliminary results of pH, EC, TDS levels that can affect human health as well as the machinery systems used for domestic and industrials purposes in the study area. Learning the variations of pH,EC and TDS levels may aid in research advancement and reduce decision-making uncertainties (Prieto-Amparán et al., 2018; Mohamed et al., 2019). For the benefit of community welfare, the study offers baseline data on water quality, which may also be used in future planning for the regional water resources, human health, and machinery system.

Description of the study area

Ambala is a rapidly expanding district in the state of Haryana that covers a total geographic area of approximately 1574 square kilometers and located between latitude 30° 10'31° 35' N and longitude 76° 30' 77° 10' E Different sites selected from different blocks are shown in Figure 1 and 2. A pronounced continental climate prevails in Ambala for most of the year where winters are considerably cold, and summers very hot. The temperatures in May and June may reach to above 46°C, while in winters can drop as low as 2°C (Paul et al., 2014). Ambala experiences both a tropical and a semi-arid climate. Thirty percent of the yearly rainfall occurs between December and February, and seventy percent occurs between July and September months with average rainfall of 47.16 mm (Ambala City Population Census 2011, Haryana) The sediments are made up of medium- and fine-grained sand particles, as well as clays, silts, and gravels in the district's western and southwest regions. The district has thin clay and sand bed inter layers and is located beneath the Ghaggar River basin. The clays are typically sticky to silty with color from brown to yellowish. Ambala, the district headquarter, is roughly 43 km from Chandigarh (U.T.) and 215 km from New Delhi, the Indian capital(Singh, N. et al., 2019). The Tangri, Beghna, and Markanda seasonal streams transverse and drain the Indo-Gangetic alluvium that covers the district (Singh, R., 2014).

MATERIALS AND METHODS

Water samples were collected from 20 different sites in different blocks of Ambala district and analyzed for pH,EC, and TDS as per standard methods (APHA2012& BIS 2012). The sampling was conducted during winter and summer seasons (2021-2022) collecting 40 samples from different locations. All water samples were collected in polypropylene bottles pre-washed with distilled water and 10% nitric acid (HNO3). Every sample was collected after running the tap for 5 minutes to avoid any major contamination from the pipes and the taps. A portion of each water sample was acidified with 98 % HNO3- during analysis. The pH of groundwater samples was determined using a portable pH meter at the sampling locations.EC of the samples was determined using the EC 372 (Systronics, India), and the TDS of the samples water was quantified using the Water Quality Analyzer PE 138 (Elico Ltd., Hyderabad, India). Using the inverse distance weighted (IDW) interpolation technique in Arc GIS 10.3 software, the maps were created for observed data. Water quality data sets are spatially represented using the geographic information system (GIS) method to create maps and perform geographical comparisons of the data.





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RESULTS AND DISCUSSION

The maps displaying the levels of pH, EC and TDS in supply water are shown in figures (Figure 3-8) covering six blocks in the Ambala district being utilized for both drinking and irrigation. Arc GIS software was used to create the maps of the collected data on supply water quality for this region.

pH variations

One of the key elements affecting water quality is pH, which decides the acidic or alkaline nature of water. (Table -1). The pH values had a mean of 7.6 and varied from 6.9 to 8.4 in winter 2021 and ranged 6.8 to 7.9 with a mean value of 7.3 in summer 2022. Values remained within the range according to WHO and BIS during both the seasons. Figure 3, shows the mapping of pH levels in different blocks of Ambala district during winter (2021) and summer (2022).

Electrical conductivity

Electrical conductivity, measures (μ S/cm) the water capacity to conduct electrical current. Water may conduct electricity because it contains ions like sodium, potassium, and chloride etc. It is an indirect measure of total dissolved salts content. The amount of ions present, and the temperature of water are crucial components of the conductivity measurements. In present study the EC values ranged between 334 and 1885 (μ S/cm) in year 2021 and 446 and 1576 (μ S/cm) in year 2022, with corresponding mean values of 625.85 and 744.05 (μ S/cm). The allowed EC value for drinking water, according to WHO (2011) standards, is 750 (μ S/cm). EC values measured during 2021-2022 were found to be higher at some areas than the regulatory limits at some sites.EC values were are out of range in year 2021-2022.

Total dissolved solid (TDS)

Soluble solids are the mineral components that are dissolved in water. Natural water typically contains < 500 mgl⁻¹ of dissolved solids, whereas water that contains greater than this amount is considered unsuitable for consumption as well as several industrial uses. Water containing a TDS content of < 300 mgL⁻¹is ideal for pulp paper, plastics, and textile dyeing. The total concentration of dissolved minerals in the water is broad indicator of the water's overall fitness for a variety of uses. Solar distillation, electro dialysis, ion exchange, and reverse osmosis methods are frequently used for removing TDS. TDS represents the amount of non-volatile chemicals in colloidal and molecularly dispersed form that are present in the water. Water contains a wide range of dissolved inorganic substances in varying concentrations because of surface and subsurface movements. It all comes down to the concentration of NaCl, which controls the conductivity and affects TDS. In 2021, TDS values ranged 200.4-1131 mgL⁻¹ with a mean value 375.51 mgL⁻¹. In 2022, TDS values ranged from 267.6 – 945.6 mgL⁻¹ with a mean value 448.21 mgL⁻¹. At some sites TDS values were out of range according to BIS and WHO Mapping of TDS levels for 2021 and 2022are shown in (Figure-7 and 8).

CONCLUSION

In order to look into the water quality for irrigation and drinking, the criteria pertaining to groundwater quality were examined. To visualise the water quality data. Arc GIS software was utilized for mapping and analysis of data. In several blocks of the Ambala district, where industrial and agricultural practices predominate, high levels of EC and TDS were detected in winter and summer months of 2021-2022. The current finding can be utilized to check the suitability of water for irrigation and domestic consumption.

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Table 1: Grid locations and abbreviations of sampling sites selected from different locks of Ambala District in Haryana

Block Name	Site Name	Site Abbreviation	Latitude	Longitude
Barara	Holi	S ₁	30.235277	77.071556
	Dosadka	S ₂	30.290048	77.113397
	Mullana	S ₃	30.275337	77.049278
	Sohana	S ₄	30.280845	76.951791
Saha	Kalpi	S ₅	30.294482	76.988794
	Saha	S ₆	30.300613	76.967186
	Tepla	S ₇	30.306392	76.946268
	Dhakola	S ₈	30.280598	76.973487
Shahzadpur	Patherari	S ₉	30.415204	77.027899
	Shahzadpur	S ₁₀	30.446484	77.033943
	Bhundmajra	S ₁₁	30.440053	77.038965





	Majra	S ₁₂	30.442322	77.032984
Naraingarh	Bara-garh	S ₁₃	30.460031	77.064972
	Badi – Basi	S ₁₄	30.470678	77.072202
	Choti-Basi	S ₁₅	30.462239	77.086309
	Chajal–Majra	S ₁₆	30.458244	77.082691
Ambala	Sullar	S ₁₇	30.316313	76.700875
	Ballana	S ₁₈	30.320678	76.728512
	Ambala Cantonment	S ₁₉	30.339599	76.830023
	Ambala City	S ₂₀	30.378180	76.776695

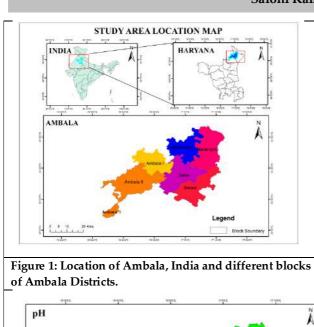
Table 2.Chemical variation of parameters analyzed in supply water Samples during 2021-2022

	2021		2022			
Sampling sites	рН	EC (μS/cm)	TDS(mgL-1)	рН	EC (μS/cm)	TDS(mgL-1)
S ₁	7.9	344	206.4	7.9	446	267.6
S ₂	7.7	334	200.4	7.8	549	329.4
S ₃	7.8	360	216	7.9	526	315.6
S ₄	7.7	345	207	7.8	546	327.6
S ₅	7.7	478	286.8	7.3	1073	643.8
S ₆	7.1	543	325.8	7.8	521	312.6
S ₇	7.2	498	298.8	7.2	519	311.4
S_8	7.6	525	315	7.7	533	319.8
S ₉	7.6	636	381.6	6.9	688	-
S ₁₀	7.5	629	377.4	7.5	614	368.4
S ₁₁	6.9	590	354	7.1	816	489.6
S ₁₂	7.2	582	349.2	7.2	605	363
S ₁₃	7.65	545	327	7.2	549	329.4
S ₁₄	7.61	754	452.4	7.3	759	455.4
S ₁₅	7.8	547	328.2	7.5	572	343.2
S ₁₆	7.5	557	334.2	7.3	759	455.4
S ₁₇	8.4	526	315.6	6.8	1027	616.2
S ₁₈	8.1	471	282.6	7.1	1059	635.4
S ₁₉	7.6	1885	1131	7	1144	686.6
S ₂₀	7.4	1368	820.8	6.9	1576	945.6
Mean	7.60	625.85	375.51	7.36	744.05	448.21
Median	7.60	544	326.4	7.3	609.5	363
S.D.	0.3466	368.2	220.9	0.356001	291.1692	179.31
Min	6.9	334	200.4	6.8	446	267.6
Max	8.4	1885	1131	7.9	1576	945.6





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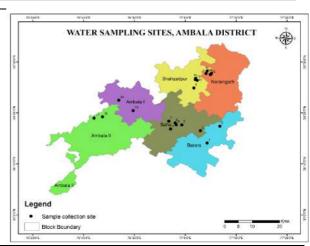
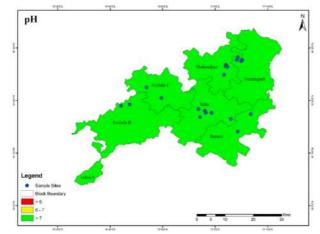


Figure 2: Water sampling sites in different blocks of Ambala.



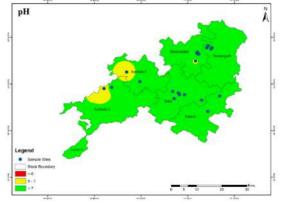
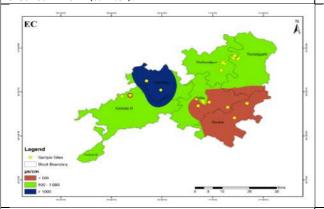


Figure 3. Mapping of pH in different blocks of Ambala district in 2021 (winter).

Figure 4. Mapping of pH in different blocks of Ambala district in 2022 (summer).



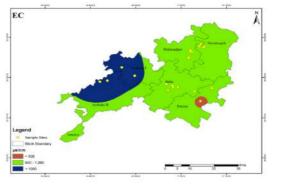


Figure -5 Mapping of EC in different blocks of Ambala district in winters 2021.

Figure -6 Mapping of EC in different blocks of Ambala district in summer 2022.





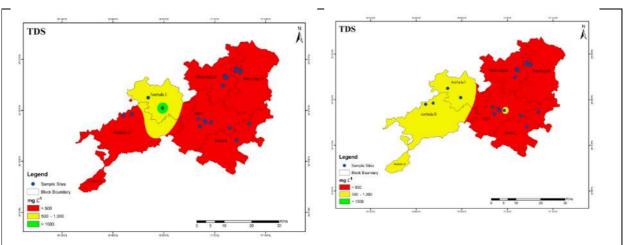


Figure-7 Mapping of TDS in different blocks of Ambala district in year 2021.

Figure 8. Mapping of TDS in different blocks of Ambala district in year 2022.





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RESEARCH ARTICLE

Fluorine-Free Hydrophobic Coating on Polyurethane Foam for Oil Spill **Absorption from Water Surface**

Lipika and Arun K. Singh*

Department of Chemistry, M. M. Engineering College, Maharishi Markandeshwar (Deemed to be University), Mullana, Ambala, Haryana 133207, India.

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*Address for Correspondence Arun K. Singh

Department of Chemistry, M. M. Engineering College, Maharishi Markandeshwar (Deemed to be University),

Mullana, Ambala, Haryana 133207, India. Email: arunkumar.singh@mmumullana.org

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ABSTRACT

In this study, polyurethane foam was functionalized by Fe₃O₄ nanoparticles and calcium stearate (CS) as a coating material to produce hydrophobic and magnetic material applicable in separating oil/water mixtures. Scanning electron microscopy (SEM), Fourier-transform infrared spectroscopy (FTIR) techniques-based analysis and water contact angle (WCA) measurement were performed to examine the surface characteristics properties of coated foam. With a water contact angle of 135°, the created foam demonstrated good hydrophobicity. The as-prepared coated foam exhibited good selective sorption capacity of oils and organic solvents from contaminated water surfaces. The prepared hydrophobic foam along with the incorporation of Fe₃O₄ nanoparticles was easily separable from treated oil/water mixtures with the use of simple conventional magnet. This study provides promising approach for the fabrication of hydrophobic and magnetic foam without use of any fluorinated coating materials.

Keywords: Hydrophobic; Oil/water mixtures, Magnetic foam, Absorption capacity

INTRODUCTION

The rapid growth in global industrialization is one of the significant reasons for the pollution in aquatic environment [1] [2]. For instance, oily contamination in aquatic body causes several adverse effects on aquatic ecosystem, leads to the water pollution and scarcity [3][4] [5]. In the recent past, several artificial surfaces with special wetting behavior such as hydrophobic and oleophilic behavior attracted wide attention for the remediation of oily contaminated water. Fabrication of such kind of super hydrophobic/ oleophilic surfaces resolves the adverse outcomes coming out from spilling of oil directly into the water and separation of oil from the water is the urgent need for the environment [6] [7].





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Currently various other techniques are used to depart oil from the oil/ water emulsion but are time taken, difficult to proceed and causes secondary pollution [8]. Foam has been used widely in the process of separating oil/water mixtures because of its three-dimensional structure[9]. Foam technology, with its elastic form, high porosity, mechanical durability, and cost-effectiveness, has earned significant interest in improving absorption capacity and streamlining oil recovery processes [10] [11] [12]. Excellent reusability, outstanding absorption capacity, and superior anti-wetting qualities are essential for the ideal absorbent material for the oil recovery process [13]. Water quality deteriorates as a result of many toxins that are present in it as a result of excessive industrial activity[14]. Contaminated oily waste water presents a serious risk to the ecosystem when it is disposed of in fresh water[15] [17]. It is currently a big problem to remove greasy materials from the water[18]. To change the hydrophilic properties of polyurethane foams into hydrophobic ones, dangerous substances as fluorinated have been employed [19][20][21].

However, fluorinated compounds required various toxic solvents for the fabrication of hydrophobic surfaces [22]. To eliminate greasy waste from water in a way that is easy to use, affordable, and environmentally beneficial, the creation of a fluorine-free unique wettable surface has gained significance [22] [23]. Oil/water separation, anti-fouling, self-cleaning, anti-microbial, anti-scaling, and anti-fogging are only a few of the applications for which the creation of superhydrophobic surfaces has become increasingly important [24] [25] [26]. In this work, non-fluorinated calcium stearate and Fe₃O₄ magnetic nanoparticles were used to generate hydrophobic surface coating on polyurethane foam with excellent magnetic behavior. Fe₃O₄ magnetic nanoparticles (NPs) were synthesized via co-precipitation approach in a basic medium [10]. Calcium stearate based coating material was used in order to provide hydrophobicity and lowering the surface energy. In addition, functionalization of Fe₃O₄ was performed in order to provide magnetically separation ability in foam from treated water [27]. The selective oily or organic solvents separation ability of the asprepared modified foam was examined in various oils-organic solvents/water mixtures.

Experimental section Materials and methods

Calcium stearate and cyclohexanol was purchased from central drug house private limited. Toluene was supplied by the Molychem. Ferric chloride hexahydrate (FeCl3.6H2O, \geq 97%), Ferrous sulfate heptahydrate (FeSO4·7H2O, \geq 98 %) and sodium hydroxide were purchased from the Loba chemie private limited. Commercial foam was purchased from local general store and rinsed with acetone and dried at room temperature for 30 min. Several oils and organic solvent were used to determine the absorption capacity such as diesel oil, engine oil, olive oil, soybean oil, n-hexane, n-heptane.

Fabrication of Magnetic polyurethane foam

The fabrication of water repellent coatings on polyurethane foam was accomplished with the help of dip coating technique. Initially, Ferrous sulfate heptahydrate and Ferric chloride hexahydrate is added in 40 ml of deionized water in molar proportion of 2:1 respectively and allowed to stirred for about 60 min. The sodium hydroxide is added to the mixture slowly until the pH of the solution reaches 11 and continued for stirring for 1 h for complete reaction. The clean small pieces of foam were done now dipped into the prepared magnetic mixture and allowed to stir for 15 min on magnetic stirrer. Finally, dipped polyurethane foam placed in hot air oven for 8 h and then dried at 60 °C for 1 h. Thus, obtained coated foam are considered as magnetic polyurethane foam.[28]

Fabrication of magnetic superhydrophobic foam

The calcium stearate coated polyurethane foam was fabricated by simple dip-coating approach by preparing the homogenous solution of 3% calcium stearate in toluene and cyclohexanol in volumetric ratio of 9:1. To decompose calcium stearate in the above mixture, heat the mixture along with continuous stirring up to 110° C. Subsequently, when the temperature decreases to 80° C, dipped the above prepared magnetic polyurethane foam into it for 5 sec and then kept for drying at 80° C for 1 h. Therefore, the fabricated foam is now regarded as magnetic hydrophobic foam.





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Characterization

The surface morphological properties of the modified polyurethane foam were examined by field emission scanning electron microscopy along with energy dispersive X-ray technique [29]. FTIR spectrometer was utilized for the determination of chemical composition on the coated and uncoated foam in the range of 400 cm⁻¹ to 4000 cm⁻¹[30]. The water repellent ability and coating durability will be examined through some scientific terms i.e., water contact angle/oil contact angle with the help of instrument known as goniometer which works on the principle of sessile liquid drop method [31].

RESULTS AND DISCUSSION

Surface morphology and chemical composition

An energy dispersive X-ray (EDS) detector and scanning electron microscopy (SEM) were utilized to investigate the elemental composition and surface morphology of coated and uncoated polyurethane foam [32]. The high-resolution FESEM spectroscopic tool has been used to investigate the surface morphology [33] and observe that the three-dimensional network-like structure of the magnetic PU foam was very porous and linked (Fig. 3a-b). High magnification SEM images made it evident that Fe₃O₄ nanoparticles were distributed randomly within the PU foam's porous structure, creating a hierarchical micro/nano structure [34]. Unmodified foam has a smoother surface than modified ones [9]. The foam have lower surface energy after being treated with calcium stearate coating solution which causes the surface to become hydrophobic[35]. The presence of Fe, Ca, and C was observed in the EDS spectra of the modified coated foam as compared to unmodified foam as shown in Fig. 3c-d. We observed two peaks at 2848 cm⁻¹, 2915cm⁻¹ and1464cm⁻¹the FTIR spectra of modified foam, corresponding to the C-H stretching vibrations and absorption peak between 1500- 1700cm⁻¹ due to symmetric and asymmetric stretching vibrations of coordinated carboxyl group in Ca(CH₃(CH₂)₁₆COO)₂[36]. Magnetic Fe₃O₄ nanoparticles have a characteristic peak that is linked to the absorption bands at around 555 cm⁻¹ (Fig.4) [9].

Evaluation of super hydrophobicity of the coated material

The water repellent ability of coated magnetic hydrophobic foam was examined by measuring water contact angle with the help of contact angle goniometer instrument via sessile drop technique at room temperature.[37] Water contact angle was measured at several diverse angle and the average value of the contact angle were taken. The water contact angle of fabricated surface was found to be.

Application in oil/water separation

In order to examined the oil/water separation capability of the coated polyurethane sponge from oil/water mixture, absorption experiments were conducted. In this experiment work, n-hexane/water, n-heptane/water, toluene/water, diesel oil/water, engine oil/water, olive oil/water, soyabean oil/water were taken (1:1 v/v) in which water was dyed with methylene blue. The oil/water mixture in 1:1 ratio was stirred vigorously for 5 min. Now, weighed amount of coated foam dipped into the oil/water mixture with continuous stirring for 5 to 7 min. and then note down the absorbed weight of the foam to calculate the absorption capability of the coated foam. In addition, the whole experimental procedure was repeated three times to calculate the mean value of absorption capacity of the coated material [38]. The oil absorption capacity was calculated by the mathematical expression. [39] [40]. The absorption capacities of the prepared hydrophobic coated foam are summarized (Table 1 and Fig. 6) with respect different oils and organic solvents.

$$K = \frac{M_1 - M_0}{M_0}$$

Where, mo= weight of foam before absorption m1= weight of foam after absorption K= absorption capacity





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CONCLUSION

A super hydrophobic/superoleophillic magnetic polyurethane (PU) foam was prepared by the functionalization with Fe₃O₄ nanoparticles and surface modification using calcium stearate based coating materials. Dip coating approach was adopted to prepare hydrophobic foam from unmodified. The developed modified foam exhibited good selective absorption capacity in the range of 13.9–50.7 g/g with respect to various oils or organic solvents from water surface. Thus results of the study inferred that as-prepared magnetic hydrophobic polyurethane foam showed excellent ability as adsorbent for oils and organic solvents from contaminated water surface.

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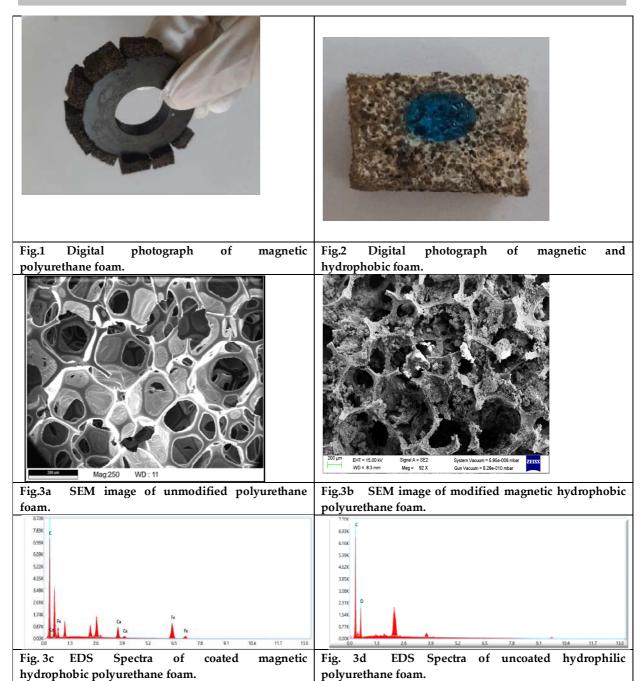
Table 1:Absorption capacities of the prepared hydrophobic coated foam are summarized with respect different oils and organic solvents.

Sr.No	Organic solver	Viscosity (cP	Density (g/cm ³	Average value (gg-
1.	n-hexane	0.31 at 25°C	0.66	13.9
2.	n-heptane	0.42 at 25°C	0.68	44.2
3.	Toluene	0.59 at 25°C	0.87	50.27
4.	Soyabean oil	50.09 at 25°C	0.91	14.7
5.	Diesel oil	3.45 at 25°C	0.82	36.95
6.	Engine oil	50 at 23°C	0.96	23.9
7.	Olive oil	84 at 20℃	0.91	34.2





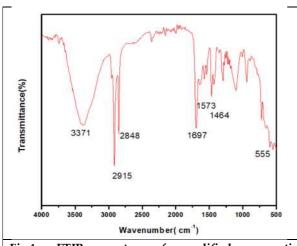
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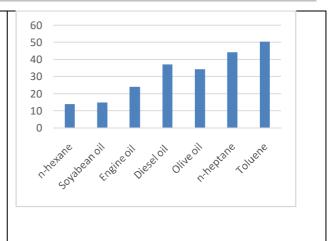


Fig.4 FTIR spectra of modified magnetic hydrophobic polyurethane foam.

Fig.5 Pictorial representation of absorption capacities of different oils/solvent.





RESEARCH ARTICLE

Gibberellic Acid Treated Seeds Improves Salinity Induced Inhibition of Germination and Seedling Growth of Thespesia populnea (L.) Soland. ex Correa, Morinda citrifolia L., and Markhamia lutea (Benth.) K.Schum

Preethi Jenifer Praticia.S 1 and Kanchana.M2*

¹Research Scholar, Department of Botany, PSGR Krishnammal College for Women, (Affiliated to Bharathiar University), Coimbatore, Tamil Nadu, India.

²Associate Professor, Department of Botany, PSGR Krishnammal College for Women, (Affiliated to Bharathiar University), Coimbatore, Tamil Nadu, India.

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*Address for Correspondence

Kanchana.M

Associate Professor, Department of Botany, PSGR Krishnammal College for Women, (Affiliated to Bharathiar University), Coimbatore, Tamil Nadu, India. Email: kanchana09psgrkcw@gmail.com



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ABSTRACT

Salinity is one of the abiotic stresses affecting crop productivity, predominantly by inhibiting seed germination. It has been demonstrated that the accumulation of soluble solutes around the seeds increases the osmotic pressure, often leading to excessive uptake of ions and impairing germination. There is some evidence to show that hormonal priming, primarily treating seeds with gibberellic acid (GA₃), is an effective strategy for regulating the germination percentage under saline stresses. Therefore, this study explored the impact of GA3 application on Thespesia populnea (L.) Soland. ex Correa, Morinda citrifolia L., Markhamia lutea (Benth.) K. Schum seeds under saline conditions. Our results demonstrate that priming seeds with 0.2 g/L GA₃ slightly enhances the germination rate from 71% to 74%. However, the germination rate decreased considerably with the rapid development of shoots and length. In particular, the mass and length of shoots were notably better compared with untreated seeds.

Keywords: Germination Percentage, Gibberellic Acid, Priming, Tolerance,





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INTRODUCTION

Salinity is a critical environmental stress affecting plant growth worldwide and one of the top ten threats to the world's soil resources [1];[2]. Previous studies have shown that salinity causes three basal issues for plants. First, excess salt in the soil decreases the osmotic potential, which leads to a reduction in water uptake by plants. Second, a higher amount of Na+ and Cl2 ion uptake distracts from the absorption of necessary minerals and imputes toxicity to plants [3]. The salinity has more adverse effects that reduce the growth, development, and yield of the crop by changing the activation of some pre-germination metabolic processes [4]. In this context, salinity affects the crop yield on over 800 million ha of arable land, which accounts for approximately 6% of the total terrestrial land and 10 million ha of agricultural land are destroyed by salt accumulation each year [5]. Seeds are more vulnerable to salinity-induced stress. Typically, seeds require more water intake during germination, which indirectly results in the accumulation of soluble solutes around the seeds, thereby increasing the osmotic pressure. In effect, the osmotic pressure change causes ion toxicity and leads to the assimilation of Na⁺ ions [6]. Hence, the ionic imbalance affects the uptake and transport of other essential ions in target cells, which stops the metabolism [7]. Many efforts have been made to improve the salinity and osmotic tolerance of the crop through selective breeding techniques [8]. However, the success rate of crop productivity is limited, and commercialization has also been less successful [9]. To mitigate the changes resulting from salt environment stress, priming, wherein seeds are treated with a chemical substance under controlled conditions, has been considered one of the simplest and most cost-effective techniques [10]; [11].

Seed germination is a complex process that includes events starting with water uptake by the (quiescent) dry seed and culminating with the elongation of the embryonic axis and the emergence of the radicle [12]. Among multiple techniques, seed priming is one of the most effective strategies by which pre-germination can be activated through physiological and chemical processes [6]; [13]. Further, seed priming will improve uniform germination by *decreasing* the imbibition time [14]. Priming treatments can be done by treating seeds with plant growth regulators, buffer solutions, and other chemical substances, e.g., gibberellic acid (GA₃) [12]; [13]. Nevertheless, a practical priming approach is accomplished by treating the seeds with GA₃, which increases the germination of fewer vigorous seeds [15]. Indeed, GA₃ as a plant hormone is known to induce different physiological responses in plants, which are well suited for stimulating and improving germination, plant growth, and photosynthetic activity [16]. Considering the tendencies of growth reduction due to salinity and the progressive efficacy of exogenous GA₃ application on different morphological, physiological, and biochemical activities, it can be described that the application of GA₃ is practical to detract salinity stress and that its effectiveness is more dynamic in salt-tolerant plants [17].

Thespesia populnea (L.) Soland. ex Correa (Malvaceae) most probably originated from the Pacific and Indian Ocean coastlines, the Asian tropics, or even both. It is currently present in coastal locations across the tropics; it is planted in coastal communities and sporadically on land [18]. One of the most significant sources of traditional remedies among Pacific Island civilizations is Morinda citrifolia L. (Rubiaceae), sometimes known as noni in the commercial sense. It grows widely throughout the Pacific. The diminutive evergreen tree or shrub is native to Australia and Southeast Asia (Indonesia), but it is now found throughout the entire tropical world. [19] and Markhamia lutea (Benth.) K. Schum (Bignoniaceae), this species natural habitat ranges from Tanzania and South Sudan to Ghana [20]. In this work, we specifically explored: (1) the germination activity and seedling growth through priming with GA3in seeds of Thespesia populnea (L.) Soland. ex Correa (Malvaceae), Morinda citrifolia L. (Rubiaceae), and Markhamia lutea (Benth.) K.Schum (Bignoniaceae), and (2) seeds were also treated with two different concentrations of NaCl, characterising their tolerance during germination and early seedling growth.





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MATERIALS AND METHODS

Seed Collection

The mature seeds of *Thespesia populnea* (L.) Soland. ex Correa (Malvaceae), *Morinda citrifolia* L. (Rubiaceae), and *Markhamia lutea* (Benth.) K. Schum (Bignoniaceae) were collected from September to December 2020 from the Western Ghats, Sadivayal area in Coimbatore district (11.0168°N, 76.9558°E). The collection site has a mean annual temperature of 26°C, ranging from 32°C in September to 19°C in December. Several hundred seeds were collected from 10 randomly chosen trees located within a 20-km radius by gently shaking the mature trees. Seeds were transported to the laboratory via road on the same day, visually inspected, and any unhealthy seeds or debris were removed. The seed surface was sterilised with a 5% sodium hypochlorite solution for 10 minutes and thoroughly washed with sterile distilled water 4–5 times. The seeds were then bench-dried for a day and subsequently stored in air-tight glass bottles under room temperature and humidity until used in the experiment. Experiments commenced within two weeks of collection.

Hormonal Priming

The seeds were primed by soaking in a 0.2 g/L GA₃ solution for 12 hours at 27°C in complete darkness, achieved by wrapping the Petri dishes with two layers of aluminium foil paper. After priming, seeds were removed and washed in running tap water, then rinsed three times with sterile distilled water. Seeds were blotted dry by placing them between filter papers. After drying, the seeds were kept in Petri dishes containing moist No. 1 Whatman filter paper at room temperature, 27°C. Control seeds (unprimed) are soaked in water for 12 hours, re-dried, and then placed in Petri dishes containing moist No. 1 Whatman filter paper in room temperature at 27°C.

A fully randomised factorial experiment consisting of three replicates of 25 seeds each from the control and primed groups was irrigated with equal volumes of NaCl solutions at two different concentrations: 50 and 100 mM. Physiological parameters such as seed germination percentage and mean germination time (MGT) were calculated by the formulae given below. The shoot and root lengths of all seedlings were measured using a transparent ruler. Shoots and roots were measured using a weighing balance. The separated shoots and roots were dried at 70°C for 12 h, and then dry weights were recorded [21]; [11].

Measurements of Germination Efficiency

Total germination (TG) was measured every day and terminated at day 14 after sowing (Kandil *et al.*, 2012). It was computed as:

 $TG\sqrt[n]{n} = \frac{n}{N} \times 100$, where *n* is the total germinated seeds, and *N* is the total seeds sown.

MGT was calculated according to Fuller et al.(2012)

 $MGT = \frac{\sum (n \times d)}{\dots}$

Where n is the number of seeds germinated on each day, d is the number of days from the beginning of the test, and N is the total number of seeds germinated at the termination of the experiment.

The vigour index (VI) was measured using the formula of [9]:

 $VI=TG\% \times Seedling length (cm)/100$

Statistical Analysis

All data collected from the experiments were subjected to two-way ANOVA using the Statistical Package for the Social Sciences (SPSS, version 20) software, and the difference between means was computed by Fisher's Post Hoc Least Significance Difference Test at P < 0.05.





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RESULTS AND DISCUSSION

Germination Percentage and Mean Germination Time of the Study Seeds

Experimental results showed that both salinity and treatment with GA3 noticeably affected the germination percentage as well as Mean Germination Time (MGT) of the three crops, Thespesia populnea, Morinda citrifolia, and Markhamia lutea at P< 0.05 (Figures 1–3). The synergistic effect of seed type, GA₃ treatment, and salt concentration also significantly affected the germination percentage as well as the MGT of Thespesia populnea, Morinda citrifolia, and Markhamia lutea at P< 0.05. Increasing the salinity concentrations in the germination medium negatively affected the germination percentage and delayed the germination time in all three species. However, the reduction was significantly more noticeable in the unprimed seeds than in the GA₃-primed ones (Figures 1-3). Notably, a high reduction was detected at the highest concentration of 100 mM NaCl. An increase in germination percentage was observed in the treated seeds at 50 mM NaCl salinity. While comparing the untreated seeds of Markhamia lutea and Morinda citrifolia with the treated ones, they were most affected due to the salinity stress. In addition, the germination rate was also significantly affected. The maximum germination rate was recorded for 0-2 g/L GA3treated seeds of Thespesia populnea at the highest salinity levels (Figures 1-3). The results of the present study are consistent with [22], who showed that plants identified the accumulation of some inorganic ions or organic compounds to balance the osmotic potential against the initial issues of salinity, which causes inhibition of water uptake by the roots due to high ion content in the soil solution. When seeds are sown in varied concentrations of salinity, the germination rate is directly proportional to the time required for germination. 0.2 g/L GA3 treated seeds show better germination at varied concentrations of salinity. GA3 can potentially increase germination percentage due to the activation of some specific genes for α -amylase mRNA transcription and shut off the defensive effects of salinity stress on germination. The progressive growth in germination percentage with GA3 might be due to the involvement of GA3 in the activation of amylases, lipases, and proteases, as well as an increase in cell wall plasticity and improved water absorption [11]. Any increase in the activity of these enzymes may result in early, vigorous germination and better crop establishment.

In addition, RNA synthesis and acid phosphatase may also increase in the cotyledons and embryonic axes of the GA3treated seeds compared with the untreated ones. The α -amylase plays a significant role in the hydrolysis of macromolecules such as proteins and starch in the aleurone cells. The stored content is converted into forms available for the development of the embryo by increasing the size and water potential through osmotic processes [23]. Besides, the signalling pathways of GA3 tend to boost seed germination through the release of seed dormancy by counteracting the activity of abscisic acid (ABA). ABA will inhibit embryonic expansion during stressful situations by weakening the endosperm and inhibiting the expansion of embryonic cells [24]. Similar effects of GA3 were observed on the germination of Sorghum bicolor [8], Ricinus communis [25], Satureja thymbra [26], Delonix regia [27], and onLathyrussativus [28].

Effect of Salinity and Priming on the Seedling Growth Performance of the Three Seeds

Seeds exposed to higher salt concentrations during germination caused stress and decreased the seedling vigour index, shoot and root length, and shoot and root fresh and dry weights of all the experimental seedlings (Figures 1–3). The salinity levels are indirectly proportional to the growth of the seedlings. It was noticed that when salinity levels increased, seedling growth decreased significantly. However, priming with GA₃ had a stimulatory effect on maximum physiological characteristics up to a certain salinity level (50 mM NaCl). An exceptionally high number of seedlings were documented at primed seedlings (75%) irrigated with salinity levels up to 50 mM NaCl for *Thespesia populnea*, followed by *Morinda citrifolia* (64%) and *Markhamia lutea* (43%) (Figures 1–3).

The seedling growth of *Thespesia populnea, Morinda citrifolia,* and *Markhamia lutea* significantly decreased when grown in high salinity. Seedling vigour declined from the control (no salt) to the 50 mM NaCl medium. The maximum seedling vigour index was observed at 50 mM NaCl (43.10 for primed and 14.12 for unprimed), and the minimum seedling vigour index was at 100 mM NaCl (3.02 for primed and 1.12 for unprimed). Figures 1–3 show reductions in





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seedlings were observed on all tested seedling traits: seedling length (SL), seed dry weight (SDW), seed fresh weight (SFW), root length (RL), root fresh weight (RFW), and root dry weight (RDW). When viewed in the context of ion uptake, high salinity in the growth medium reduces the water uptake capacity of the seedlings, which causes a reduction in growth rate. The mechanism by which salt tolerance affects the growth of a plant depends on the time scale, and the plant is exposed to a high salt environment. Na⁺ and Clions can enter the cells and influence the metabolic activities in the cytosol and cell membranes via direct toxic effects [29]. [30] concluded that seed priming with GA₃ in *Calendula officinalis* L. and *Foeniculum vulgare* Mill. accelerated metabolic activity and increased radicle and plumule weight, especially under salt stress.

The roots of all the tree seeds were affected first due to direct contact with the saline growth medium, then the shoots of all tree seeds (Figures 1–3). This is because the roots were under osmotic salt pressure at the rhizopsheric region of the seedlings, which directly corroded the roots. The osmotic regulation, together with the toxic effects of Na⁺ and Clions, decreases water uptake and causes an imbalance of essential nutrients at the time of seed germination (31). GA₃ caused a significant reduction in Na⁺ content as well as decreased the range of other ions [32]. Similar results were also observed in stressed maize plants treated with GA₃, showing a significant reduction in the accumulation of fewer Na⁺ ions [33]. Most toxic effects of NaCl can be attributed to Na⁺ toxicity, resulting in the dormancy of seeds and delayed germination. In addition, K⁺ ions also play a vital role in stomatal regulation and preventing water loss and necrosis. Hence, it is essential to nullify the toxicity of the Na⁺ ion, and nevertheless, it increases the uptake of K⁺ ions to improve cellular function [34]

CONCLUSION

This study provided valuable information about *Thespesia populnea, Morinda citrifolia,* and *Markhamia lutea* seeds under salt stress and pretreatments with a growth hormone (GA₃). Seed priming with GA₃ significantly improved seed germination and seedling growth and alleviated the inhibitory effect of salt stress on germination and seedling growth of the investigated tree plants. Decreases in seed germination percentage and seedling growth with increasing salt stress were more apparent in the unprimed seeds than in the primed seeds. Furthermore, the roots of the studied tree plants were more affected than the shoots. Pre-treatment with GA₃ increased total seed germination percentage, decreased MGT, and increased seedling growth performances. In conclusion, this study showed the effectiveness of applying seed priming techniques in salt-stressed environments to reduce the harmful effects of salinity stress on seed germination and the early growth of the seedlings *in vitro*. Nevertheless, further research is necessary to see the performance of the three primed (GA₃) tree plants on vegetative growth and yield under field conditions.

Conflict of interest

The authors declares that there is no of conflict of interest

ACKNOWLEDGEMENT

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Table:1 Analysis of variance for morphological growth parameters of *Thespesia populnea*, *Morinda citrifolia*, and *Markhamia luteas*eeds primed with gibberellic acid and germinated using various concentrations of NaCl solutions

Source of Variation	Total Germination (%)	Mean Germination Time (days)	Seedling Vigour (SVI)	Shoot Length (cm)	Root Length (cm)	Shoot Fresh Weight (g)	Shoot Dry Weight(g)	Root Fresh Weight (g)	Root Dry Weight (g)
Treatment	8572.33*	50.94*	464*	18.08*	14.02*	4.35*	2.3*	2.49*	1.38*
seed type	2056.48*	55.40*	50*	9.75*	6.75*	5.38*	1.1*	5.41*	2.11*
Salt conc.	6097.71*	53.70*	14*	4.43*	4.3*	1.23*	1.3*	3.29*	3.01*
Treatment × seed type	354.73*	0.29*	1.3*	1.73*	5.4*	3.23*	0.52*	1.73*	4.01*
Treatment × salt conc.	1094.97*	0.90*	4.5*	2.13*	6.7*	6.11*	4.35*	2.49*	2.38*
seed type× salt conc. Treatment × seed type ×salt conc	363.09* 706.73*	0.42* 0.68*	1.2* 8.2*	4.73* 5.27*	8.1* 7.9*	3.58* 2.11*	3.72* 0.91*	1.39* 0.58*	3.59* 0.27*
Error	14.73	0.8	1.82	3.72	0.054	0.0015	0.067	0.269	0.0084

^{*}Significantly different at *P*< 0.05.

Table :2 Correlation analysis between germination and morphological growth parameters of *Thespesia* populnea, Morinda citrifolia, and Markhamia lutea seeds primed with gibberellic acid and germinated using various concentrations of NaCl solutions

	using various concentrations of the crossations								
	SL	RL	SFW	SDW	RFW	RDW	GR	MGT	SVI
SL	1	0.81	0.85	-0.75	-0.73	0.95	0.98	-0.59	0.96
RL		1	0.87	-0.99	-0.56	0.49	0.95	-0.54	0.72
SFW			1	0.93	0.72	0.72	0.41	-0.47	0.39
SDW				1	0.86	-0.76	0.75	-0.67	0.81
RFW					1	0.78	0.67	-0.73	0.77
RDW						1	0.49	-0.61	0.49
GR							1	0.72	0.73
MGT								1	0.59
SVI									1

SL: length of shoot, RL: length of root, SFW: shoot fresh weight, SDW: shoot dry weight, RFW: Root fresh weight, RDW: Root dry weight MGT: Mean germination time, GR: Germination rate, SVI: Seedling vigour index.





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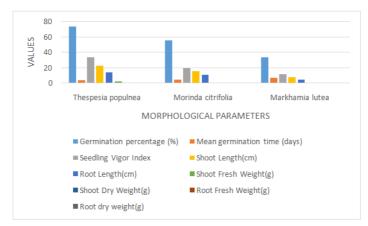


Fig :1 Effect of seed priming with gibberellic acid for morphological growth parameters of *Thespesia populnea*, *Morinda citrifolia*, and *Markhamia lutea*seeds primed after germinated under salinity stress (50mMNaCl concentration)

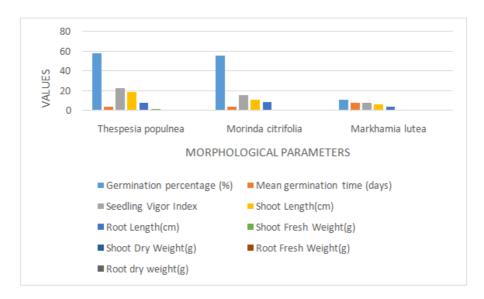


Fig: 2 Effect of seed priming with gibberellic acid for morphological growth parameters of *Thespesia* populnea, Morinda citrifolia, and Markhamia luteaseeds primed after germinated under salinity stress (100mMNaCl concentration)





Preethi Jenifer Praticia and Kanchana

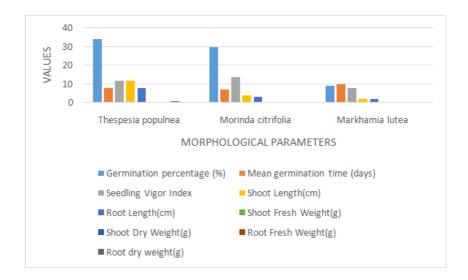


Fig: 3 Morphological growth parameters of *Thespesia populnea*, *Morinda citrifolia*, and *Markhamia lutea*unprimed seeds (without GA₃) after germinated under salinity stress (50mM and100mMNaCl concentration)





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RESEARCH ARTICLE

Method Development and **Validation** of Chloramphenicol Commercially Available Milk by LC-MS/MS

Syed Rihana^{1*}, Vidyadhara Suryadevara² and Venkata Rao Vutla³

¹Assistant Professor, Department of Pharmaceutical Analysis, Chebrolu Hanumaiah Institute of Pharmaceutical Sciences, Guntur, Andhra Pradesh, India.

²Principal and Professor, Department of Pharmaceutics, Chebrolu Hanumaiah Institute of Pharmaceutical Sciences, Guntur, Andhra Pradesh, India.

³Professor, Department of Pharmaceutics, Chebrolu Hanumaiah Institute of Pharmaceutical Sciences, Guntur, Andhra Pradesh, India.

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*Address for Correspondence

Syed Rihana

Assistant Professor,

Department of Pharmaceutical Analysis,

Chebrolu Hanumaiah Institute of Pharmaceutical Sciences,

Guntur, Andhra Pradesh, India. Email: rihanasyed687@gmail.com

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ABSTRACT

A quantitative method for the determination of chloramphenicol in milk samples was developed based on the (quick, easy, cheap, effective, rugged, and safe) approach for lc-ms/ms chromatographic method for determination of chloramphenicol in milk using 0.1% formic acid in hplc grade water: acetonitrile as mobile phase at 0.6 ml/min with agilent eclipse plus c18, (5 μm ,4.6×150 mm) as column using triple quadrapole mass spectroscopy was developed. The developed method was validated and parameters were found to be within limits The validation experiments demonistrated system precision, system suitability, accuracy, linearity of experimental parameters. The retention time of system suitability are given chromatography-tandem mass spectrometry (LC-MS/MS). Homogenized milk samples were extracted with acetonitrile. Validation results demonstrated that this method fulfills criteria for the determination of chloramphenicol in milk. The present study was carried out to validate a simple ,precise and accurate lc-ms/ms chromatographic method for determination of chloramphenicol in milk using 0.1% formic acid in hplc grade water: acetonitrile as mobile phase at 0.6 ml/min with agilent eclipse plus c18 ,(5 µm ,4.6×150 mm) as column using triple quadrapole mass spectroscopy was developed. The developed method was validated and parameters were found to be within limits

Keywords: Chloramphenicol; LC-MS/MS; Milk; Liquid-Liquid Extraction (LLE)





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INTRODUCTION

Antibiotics, also called anti-bacterials, are a type of antimicrobial drug used in the treatment and prevention of bacterial infections. They may either kill or inhibit the growth of bacteria. A limited number of antibiotics also possess antiprotozoal activity. Antibiotics are used in different food commodities to promote the growth rate and efficacy, for a lactation property, etc. Antibiotics in milk samples, which might pose a potential risk for human health due to their sub acute and chronic toxicity. Antibiotics in milk yields to severe acute and long term health effects like cancer, leukemia, grey baby syndrome in infants, Aplastic anemia, Bone marrow suppression, Neurotoxic reactions, Hypersensitivity reactions. When we consume the milk that contains more amount of antibiotics it leads to acute and chronic health problems. So, antibiotic usage should be reduced. Chloramphenicol is a broad-spectrum antibiotic suitable for the treatment of variety of infectious organisms (Bayo et al. 1994). The compound has been bannedin several countries, including the European Union, for treatment of food-producing animals. The European Commission defined aminimum required performance limit for chloramphenicol in food of animal origin at a level of 0.3 µg kg-1. However, data from p a s t years from Rapid Alert System for Food and Feed indicated contamination of incidents in various matrices with chloramphenicol (456 cases) in which 49 were in dairy products. Different analytical methods have been developed for the determination of chloramphenicol in milk using ELISA (Ferguson et al. 2005; Shi et al. 2010; Tao et al. 2012; van der Water and Haagsma 1991), gas chromatography (Kijak 1994.; Pfenning et al. 1998), and liquid chromatography(Ali et al. 2009; Bayo et al. 1994; Berendsen et al. 2013; Guy et al. 2004; Rezende, Fleury Filho, and Rocha 2012; Rodziewicz and Zawadzka 2008; Shi et al. 2010). Most of these methods are based on LLE (Dubourg et al. 1987; Perez et al. 2002; Petz and Lebens 1983; Ramsey et al. 1998; Rezende et al. 2012; Wal, Peleran, and Bortes 1980), solid phase extraction (SPE) (Guy et al. 2004; Rejtharová and Rejthar 2009), or combination of both (Sniegocki, Posyniak, and Z'mudzki 2007). In recent years, a fast and inexpensive approach for the determination of pesticide residues in fruit and vegetable samples, named QuEChERS was developed by Anastassiades et al. (2003). However, few studies have been reported for the determination of different class of veterinary drugs in samples of animal originsuch bovine milk and liver (Kinsella et al. 2009), and shrimp (Villar-Pulido et al. 2011). Only one paper reported the determination of forbidden compounds such as chloramphenicol in honey (Pan et al. 2006). The goal of this work was to develop a simple and low-cost procedure LLE methods for the determination of chloramphenicol in milk.

EXPERIMENTAL

Reagents and Materials

Acetonitrile was obtained from Merck (HPLC)graded.. Ammonium formate obtained from Merck which is Ar grade. Water which is obtained from rankem which is HPLC graded. Formic acid was obtained from Merck (Emsure). All reagents were of analytical grade or higher The liquid chromatographic system consists of the following components: the hplc system shimadzu consisting of quaternary pump, degasser, auto sampler, thermo stated column oven compartment, isocratic elution with mass detector, computer & printer and mass spectrometer consists of the following components: the mass system, mass detector, roughing pump, syringe pump and a gas generator. Chromatographic analysis was performed using lab solutions software on a column agilent eclipse plus $C18, 5 \mu m$, $4.6 \times 150 \mu m$.

Certified Reference Standards

During method validation, a blank matrix will be spiked with the analyte of interest using solutions of reference standard. It is important that the quality of the reference standard is ensured, as the quality (purity) may affect the outcome of the analysis, and therefore the outcome of the study data. Therefore the reference standards used for the analytical validation and analysis should be obtained from an authentic and traceable source. Suitable reference standards, include certified standards such as chloramphenical d5(sigma Aldrich,Dr.ehrenstorfer), commercial available standards. Suitability of the reference standard should be scientifically justified. They should be stored in frost free refrigerator below -10oc in order to prevent undesirable reactions.





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COLLECTION OF SAMPLES

Different brands of milk samples were collected from dairy in hyderabad . The milk samples were homogenized, extracted and analyzed as per the sample preparation procedure. On the basis of analysis the compound were not detected and their results are reported.

SAMPLE PREPARATION

Liquid-Liquid Extraction (LLE)

Take 2 ml of homogenized sample were weighed and transferred in to 50 ml centrifuge tube

To this add internal standard to all samples, spiked samples too then extract with ethyle acetate. For separation of compound. Then vortex for 5 minutes & centrifuge for 10 minutes. Transfer the 1 ml of supernated liquid into ria vials then it undergone to the nitrogen stream turbo evaporator for 15 minutes. It was reconstricted with diluents then clean up with hexane. Finally centrifuge it for 5 minutes and transfer the bottom layer of the sample in to instrument vails then inject the sample to lc-ms/ms Selection of initial conditions for method development Determination of

solubility of chloramphenicol

Chloramphenicol is freely soluble in methanol and chloroform.

Selection of stationary phase and mobile phase

For selection of stationary phase, two different columns C18 and C8 were used in order to get optimum results and various solvent systems such as acetonitrile, buffers and methanol. buffer were used in various ratios.

Preparation of calibration curve standards

Prepare calibration curve standards in samples from the above working standard solution as shown in below table and process the samples.

EXTRACTION PROCEDURE

Take 2 ml of homogenized sample in 50 ml centrifuge tube. Extract the sample with 10 ml ethyl acetate vortex for 5 minutes, then centrifuge at 4000 rpm. Transfer about 1 ml of supernated liquid & evaporate under nitrogen stream. By using turbo evaporater Reconstritute with 0.2 ml of diluent then vortex for 5 minutes, Clean up the sample by adding 1 ml of hexane. Transfer the bottom layer in to auto sampler vial & inject in to lc-ms/ms

Preparation of QC stock dilutions

Prepare QC stock dilution standards in samples from the above working standard solution as shown in below table and process the samples

Preparation of QC stock solutions

Preparation of 0.3ppb QC stock solution: Weigh 10gm of sample and spiked with 0.060ml of 10 ppb mix into sample and followed the sample preparation procedure.

Preparation of 0.5ppb QC stock solution

Weigh 10gm of sample and spiked with 0.100ml of 10.0 ppb mix into sample and followed the sample preparation procedure.

Preparation of 1ppb QC stock solution

Weigh 10gm of sample and spiked with 0.200ml of 10.0 ppb mix into sample and followed the sample preparation procedure.





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RESULTS AND DISCUSSION

DATA PROCESSING

Acquired chromatograms using the computer based chemstation software, process the data by peak area method. The concentration of the unknown is calculated by using regression analysis.

y = mx + c

Where, x = concentration of analyte m = slope of calibration curve

y = peak area of analyte

c = y-axis intercept of the calibration curve Calculation

Concentration $\mu g/kg = \text{concentration found } x \text{ volume of extract made } \mu p/\text{wt.of sample taken for analysis.}$

Use a regression equation with an appropriate weighting factor for determining the detector response/ concentration relationship. Include standard a and standard i in preparation of calibration curve The representative calibration curve for regression analysis is illustrated. Linearity was established using matrix-matched calibration curves. five calibration curves were prepared at levels of Matrix blank, 0.1ppb, 0.3ppb, 0.5ppb, 1ppb and 2ppb, in milk.

These curves were prepared and plot a graph through software against the concentration of the each Analyte and its response. The Correlation Coefficient (R2) \geq 0.99.

SYSTEM SUITABILITY

System suitability shall perform by injecting six replicates of known concentration of 1.00 ppb aqueous standard solution of Chloramphenicol & Chloramphenicol d5 .% RSD of area was >20% and retention time was within ± 2 min. Mean= no. of observations /total no. of runs .

Standard deviation= _____% cv= std.dev/mean volume*10.

PRECISION

The precision of the analytical method describes the closeness of repeated individual measures of analyte. Precision is expressed as the coefficient of variation (CV). The statistical method for estimation of the precision should be predefined and calculated according standard practice. Method precision was performed by injecting six spiked samples at 0.3 ppb,0.5 ppb and 1 ppb. % RSD of area \leq 20% and % Accuracy (Recovery) was injecting the spiked sample at 0.3ppb, 0.5ppb and 1ppb levels. Acceptance criteria in between 80-120%.

Formula: Precision=standard deviation /mean volume* 100

Accuracy

The accuracy of an analytical method described the closeness of the determined value obtained by the method to the true concentration of the analyte (expressed in percentage). Accuracy should be assessed on samples spiked with known amounts of the analyte, to the quality control samples (QC samples). During method validation accuracy should be determined by replicate analysis using a minimum of 6 determinations at a minimum of 3 concentration levels which are covering the calibration curve range. The QC samples are analyzed against the calibration curve, and the obtained concentrations are compared with the nominal value . Method accuracy was performed by injecting six spiked samples at 0.3,0.5,1.00ppb. % RSD of area $\leq 20\%$ and % Accuracy (Recovery) was injecting the spiked sample at 0.3ppb, 0.5ppb and 1ppb levels. Acceptance criteria in between 80-120%.

Accuracy=mean volume /set volume*100

BATCH ANALYSIS

Batch analysis was performed by collecting different brands of milk samples from Diary forms and they are analyzed by using the LLE method and the results are tabulated below. The results are expressed in ppb level i.e; ($\mu g/kg$) The present study was carried out to validate a simple ,precise and accurate lc-ms/ms chromatographic method for determination of chloramphenicol in milk using 0.1% formic acid in hplc grade water : acetonitrile as mobile phase at 0.6 ml/min with agilent eclipse plus c18 ,(5 μ m,4.6×150 mm) as column using triple quadrapole mass spectroscopy was developed. The developed method was validated and parameters were found to be within limits. The validation experiments demonistrated system precision , system suitability , accuracy, linearity of experimental





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parameters. The retention time of system suitability are givin in table -5.12, precision results are found in table -5.14 for 0.3 ppb,0.5 ppb and 1 ppb. the chromatograms for the precision study were shown.

CONCLUSIONS

A simple, sensitive, and high-throughput method for chloramphenicol determination in milk samples was developed and validated. The developed LLS technique for the determination of chloramphenicol allows extraction and clean-up to be carried out in simple steps, without additional purification of the extracts. Good values for validation parameters such as linearity, recovery. The method was developed and validated for the estimation of chloramphenicol in different brands of milk by lc-ms/ms. Milk samples were collected from Dairy forms in hyderabad and were analyzed. On the basis of analysis I conclude that chloramphenicol were not detected in milk so it is beyond the limit in milk. so we are in safe zone. Antibiotics are used in different food commodities to promote the growth rate and efficacy, for a lactation property, etc. Antibiotics in milk yields to severe acute and long term health effects like cancer, leukemia, grey baby syndrome in infants, Aplastic anemia, Bone marrow suppression Neurotoxic reactions, Hypersensitivity reactions. When we consume the milk that contain more amount of antibiotics it leads to acute and chronic health problems. So antibiotic usage should be reduced.

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Table 1.1:Preparation of diluents working standard solution

Stock		Volume of stock	Volume of methanol	Final	Final
conc.(ppm)		standard(ml)	(ml)	volume(ml)	conc.(ppb)
10.00	0	0.100	0.900	1.000	1000.00
1.0		0.100	9.900	10.000	10.00
0.01		0.100	0.900	1.000	1.00

Table 1.2:Preparation of calibration curve standards

Working standard Conc.(ppb)	Volume of standard (ml)	Sample weight(gm)	Final conc.(ppb)	Level
10.00	0.400	2.0	2.00	Cc-5
10.00	0.200	2.0	1.00	Cc-4
10.00	0.100	2.0	0.50	Cc-3
10.00	0.060	2.0	0.30	Cc-2
1.00	0.200	2.0	0.10	Cc-1
0.00	0.00	2.0	0.00	Blank

Table 1.3:Preparation of qc stock dilutions

_ rable risk repairmen or desiren				
Working standard Conc.(ppb)	Volume of standard(ml)	Sample weight(gm)	Final conc.(ppb)	Level
10.00	0.200	2.0	1.00	Qc- 1.0ppb
10.00	0.100	2.0	0.50	Qc- 0.5ppb
10.00	0.060	2.0	0.30	Qc-





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0.3ppb

Table1.4:Mrm parameters

Table1.4:Mrm parame	eters							
Compound	Precursorion	Production	D well time	Q1Prebase(v)	CE	Q3	RT	Polarity
name	ms 1 response	ms 2	(m.sec)			Prebas		
		response				e		
Chloramphenicol		152.05		11	15	16		Negative
				15	27	11		
	321						2.5	
	021	121						
			100	20	10	11		
			100					
		257						
Chloramphenicol		157		15	16	15		Negative
D5								
				15				
	326	2/2/05				45	2.5	
	020	262.05	400		11	17		
			100					
				11	28	12		
		126						
		120						
		1						

Table1.5:Linearity For Correlation Coefficient

Compound name	R-value	R-2value
Chlorampenicol-01	0.998	0.997
Chlorampenicol-02	0.998	0.997
Chlorampenicol-03	0.998	0.997

Table 1.6:Linearity for the ion of 321>152

Linearity	Area	Actual concentration(ppb)	Obtained concentration(ppb)	Accuracy(%)
Cc1	26159	0.1	0.090	90.50
Cc2	78530	0.3	0.315	104.90
Cc3	118354	0.5	0.542	108.40
Cc4	213877	1	0.971	97.1
Cc5	406442	2	1.981	99.1

Table 1.7:Linearityfortheionof321>121.10

Linearity	Area	Actual concentration(ppb)	Obtained concentration(ppb)	Accuracy(%)
Cc1	8136	0.1	0.089	89.5
Cc2	26742	0.3	0.320	106.70
Cc3	39766	0.5	0.537	107.50





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Cc4	73195	1	0.974	94.4
Cc5	139205	2	1.979	99.0

Table 1.8: Chloramphenicol d5(istd area)326.00>157.10

Linearity	Area
Cc1	43942
Cc2	41903
Cc3	37333
Cc4	38070
Cc5	35721

Table1.9: System Suitability RT

Compound	Run-	Run-	Run-	Run-	Run	Run		Stde	
name	01	02	03	04	-05	-06	Mean	v	%cv
Chlorampenico 1-01	2.468	2.471	2.469	2.469	2.47	2.471	2.47	0.00	0.05
Chlorampenico 1-02	2.466	2.468	2.468	2.469	2.468	2.469	2.47	0.00	0.04
Chlorampenico l-d5(ISTD)	2.444	2.446	2.446	2.445	2.447	2.447	2.45	0.00	0.05

Table1.10: System Suitability Area

1 4 2 10 2 12 0 1 0 y 3 1 0 2 2 1 0 4 2 1 4 2									
Compound name	Run-01	Run- 02	Run -03	Run- 04	Run- 05	Run- 06	Mean	Stdev	%cv
Chloramphenicol	129120	13742 5	1302 84	1357 98	14104 9	14240 9	136014. 1	5452.4	4.01
Chloramphenicold 5	192503	20170 1	1864 62	1954 38	20240 5	20594 3	197408. 6	7266.3	3.68

Table 1.11:PrecisionAt 0.3ppb

Compound Name	RUN -01	RUN -02	RUN -03	RUN -04	RUN -05	RUN -06	Mean	Stdev	%C V
Chlorampenico 1-01	71515	64978	70498	67426	67922	68186	68420.8 3	2325.3 1	3.40
Chlorampenico 1-02	23728	23823	23628	22716	23365	23826	23514.3 3	426.76	1.81
Chlorampenico l-d5(ISTD)	36664	36015	37608	36908	35007	37909	36685.1 7	1064.4 1	2.90

Table1.12:Precision at 0.5ppb

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Compound Name	RUN- 01	RUN- 02	RUN- 03	RUN- 04	RUN- 05	RUN- 06	Mean	Stdev	%CV	





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Chlorampenicol-01	95819	83493	82479	89550	86408	89536	87880.83	4881.44	5.55
Chlorampenicol-02	33037	28350	26599	29346	30279	31321	29822.00	2261.56	7.58
Chlorampenicol-									
d5(ISTD)	34430	31036	28798	30334	31871	33005	31579.00	1991.27	6.31

Table 1.13:Precisionat 1 ppb

			RU	RU	RU	RU			0/ C
Compound	RUN	RUN-	N-	N-	N-	N-	Mean	Stde	%C V
Name	-01	02	03	04	05	06	Mean	v	V
Chlorampenicol-01	1559	1604	1568	151250	156826	150565	155311.	3752.	2.42
Chloralitperficor-or	64	52	10	131230	150020	130303	17	73	2,12
Chlorampenicol-02	5529	543	532	50339	52349	51988	52930.8	1767.	3.34
Chioranipenicoi-02	5	33	81	30339	32349	31900	3	66	3.34
Chloramphenic									
olD5(ISTD)	2984	298	289	27651	31003	28714	29329.3	1149.	3.92
	0	01	67	27001	31003	20/14	3	88	5.72

Table 1.14:Accuracy At0.3ppb

	RUN-	RUN	RUN	RUN	RUN	RUN		
Compound	01	-02	-03	-04	-05	-06	Mean	Accuracy
Name								
Chlorampenicol-	0.328	0.302	0.315	0.306	0.326	0.302	0.31	104.39
01								
Chlorampenicol-	0.325	0.332	0.315	0.309	0.335	0.315	0.32	107.28
02								
Chlorampenicol-	0.328	0.322	0.327	0.306	0.331	0.301	0.32	106.39
03								

Table1.15: Accuracy At 0.5ppb

Compound Name	RUN- 01	RUN- 02	RUN- 03	RUN- 04	RUN- 05	RUN- 06	Mean	Accuracy
Chlorampenicol- 01	0.474	0.458	0.488	0.504	0.462	0.462	0.47	94.93
Chlorampenicol- 02	0.484	0.46	0.465	0.488	0.479	0.478	0.48	95.13
Chlorampenicol- 03	0.474	0.462	0.469	0.501	0.469	0.478	0.48	95.10

Table 1.16:Accuracy At1 ppb

Tubic 1.10.11cculu	cy min pp	,						
Compound Name	RUN- 01	RUN- 02	RUN- 03	RUN- 04	RUN- 05	RUN- 06	Mean	Accuracy
Chlorampenicol- 01	0.903	0.93	0.935	0.945	0.873	0.906	0.92	91.53



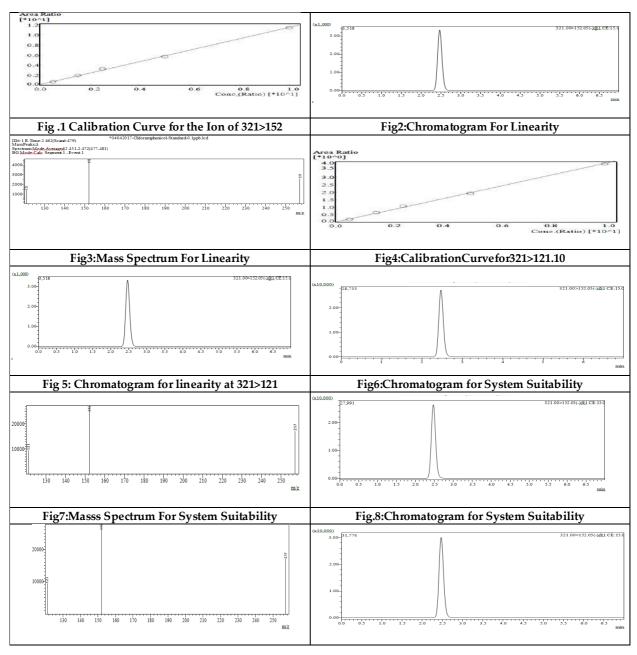


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Chlorampenicol- 02	0.939	0.923	0.932	0.922	0.855	0.917	0.91	91.47			
Chlorampenicol- 03	0.892	0.898	0.915	0.964	0.882	0.899	0.91	90.83			

Table1.17:BATCH ANALYSIS REPORT

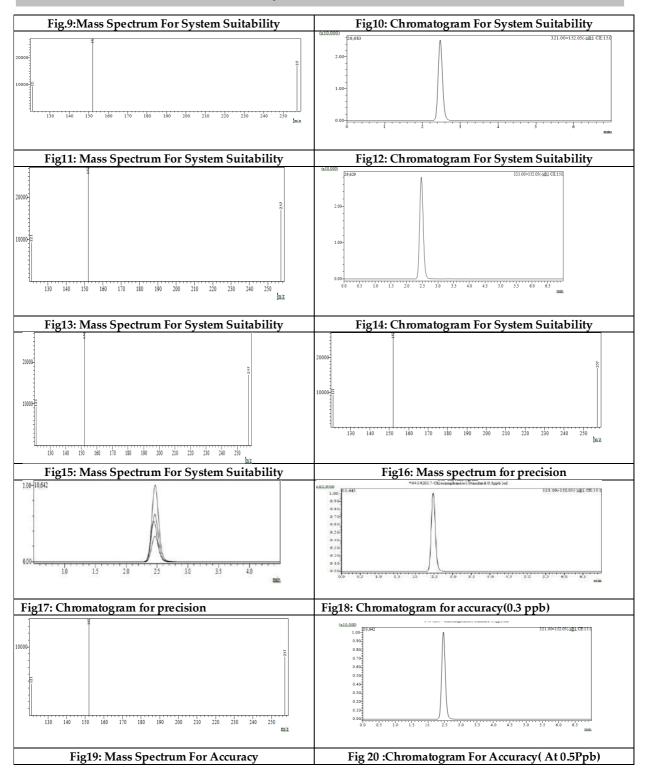
Compound name	Sample-01	Sample-02	Sample-03	Sample-04	Sample-05
Chloramphenicol	BLQ	BLQ	BLQ	BLQ	BLQ





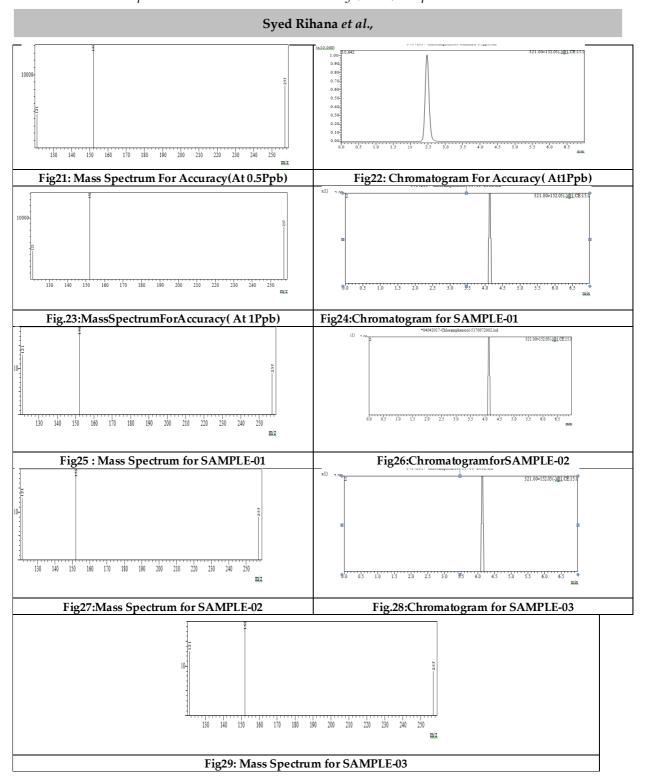


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RESEARCH ARTICLE

Alfa-Tocopherol **Enhance** Foliar-applied and Brassinolide Tolerance, Growth Improvement, and Pigment Composition in Vigna radiata L. (Co-6) Variety Grown under NaCl Stress

Aamir Abdullah¹ and R. Somasundaram^{2*}

¹Ph.D, Research Scholar, Stress Physiology Lab, Department of Botany, Annamalai University, Annamalai Nagar, Chidambaram, Tamil Nadu, India.

²Professor Stress Physiology Lab, Department of Botany, Annamalai University, Annamalai Nagar, Chidambaram, Tamil Nadu, India.

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*Address for Correspondence

R. Somasundaram

Professor Stress Physiology Lab,

Department of Botany,

Annamalai University,

Annamalai Nagar, Chidambaram, Tamil Nadu, India.

Email: botanysundaram@gmail.com



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ABSTRACT

Soil salinity is a key environmental constraint that reduces the growth development and production of essential crops globally and other abiotic stresses. Therefore, certain strategies, such as the use of plant growth regulators might be useful ways to ameliorate the adverse consequences of NaCl-induced stress on agricultural crops. In this view, the present Experimental pot culture study was conducted to determine the efficacy of foliar application of Alfa-Tocopherol (200 mg/L) Brassinolide (4 mg/L), (Alfa-Toc + NaCl), and (BL + NaCl) on (Vigna radiata L. Co-6 variety), under NaCl (80mM) concentration. however, NaCl was applied by soil drenching. The plant samples were collected after the 25th, 35th, and 45th, Days after sowing (DAS), to investigate the different physiological parameters such as root length, stem length, biomass, and pigment composition, in all applied treatments. Salt stress reduces plant growth pigment composition and biomass, Application of Alpha-Tocopherol and Brassinolide on leaves mitigates the negative effects of salt stress supporting the recovery of physiological aspects and improving pigment composition. Overall, in Vigna radiataL. (Co-6 var), Alfa-Toc and BL foliar spray mitigated the negative effects caused by NaCl stress, and it can be concluded that these plant growth regulators induce the tolerance to stress in the plants.

Keywords: Alfa-tocopherol, brassinolide, salt stress, pigment composition, NaCl.





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INTRODUCTION

Salinity is the main environmental factor affecting plant production and development. [1] In the modern world, soil salinity has emerged as a significant issue for agriculture. [2]. Approximately 800 million hectares of land globally are impacted by salinity (397 million hectares) or sodicity (434 million hectares).[3.4]. Almost all of the biochemical and physiological processes that occur in plants are impacted by the intricate mechanism of salt stress. [5] The impact of salt stress arises from the reduced osmotic potential in the soil solution, leading to water scarcity. This stress triggers a domino effect, causing imbalances in nutrients, specific ion impacts, or a blend of these factors.[6]. Salt stress may result in membrane disorder, toxic metabolite production, photosynthetic reduction, ROS production, and reduced nutrition acquisition, all of these may result to cell and plant death. [7,8]. Stress due to salinity raises Na+ and Cl- concentrations in the root, stem, and leaves, while the leaves accumulate the most ions, followed by the stem and root, demonstrating a positive association between Na+ and Cl concentration[9]. With the increase in salinity stress, a linear decrease in the levels of total chlorophyll, chlorophyll a, chlorophyll b, carotenoids, and xanthophylls as well as the intensity of chlorophyll fluorescence was found in Vigna radiataL. [10]High salt levels in the soil disrupt water balance in plants, lowering leaf water potential and affecting crucial plant functions like growth, nutrient uptake, and photosynthesis.[11]. salinity causes excessive accumulation of reactive oxygen species (ROS) such as superoxide (O2*-), hydrogen peroxide (H2O2), hydroxyl radical (OH*), and singlet oxygen (1O2) which can cause peroxidation of lipids, oxidation of protein, inactivation of enzymes, DNA damage, and interact with other vital constituents of plant cells. [12]. In south and southeast Asia, mungbean Vigna radiata L. is one of the major pulse crops. Approximately 90% of the world's production occurs in South Asia, with India being the primary producer. [13] Mungbean seeds are a good source of dietary protein for humans including marginal people, and mostly vegetarian people, they are rich sources of vitamins and minerals [14,15].

Salinity affects the mungbean plants showing a significant reduction in whole plant growth, biomass, and variation in leaf color, and photosynthesis, reducing nodule respiration, nitrogen fixation, and root hair formation[16]. Alphatocopherol acts as a crucial plant growth regulator, shielding photosystems from photo-inhibition and safeguarding chloroplast membrane lipids during salinity stress. [17]. α -Toc is effective in eliminating free radicals and preventing membrane lipid peroxidation in soybean. [18]. Using alpha-tocopherol externally has proven highly effective in boosting growth and yield-related characteristics in *Calendula officinalis*. [19]. Alpha-tocopherol application externally to *Helianthus annuus*, regulated protein content and sugar levels, ultimately enhancing yield. It also led to improved chlorophyll content, growth, and development in *Vigna radiata* L. [20,21]. Brassinolide is an important plant growth regulator enhancing the growth development, photosynthesis, photomorphogenesis, chlorophyll, and pigment composition in plants. [22]. Brassinolide acts as a guardian, easing the detrimental effects of salt stress on plants by safeguarding stomatal conductance, maintaining membrane integrity, preserving leaf hydration, and balancing the ionic composition. [23,24]. Application of Brassinolide applied 1mg/L^{-1} contributed significantly in improving the floral characteristics, highest average No. of racemes, and fertility percentage in *Vigna radiata* L.[25].

MATERIALS AND METHODS

Seed collection and Chemical reagents.

Vigna radiata L. (Green gram seeds Co-6 variety) were purchased from Tamil Nadu Agricultural University Coimbatore, Tamil Nadu, India. Chemical regulators Alfa-tocopherol and Brassinolide and analytical reagent NaCl were purchased from Sisco Research Laboratories [SRL]- Chennai-600117.

Experimental design.

The Experimental study was conducted within the Botanical Garden of Annamalai University, situated in Chidambaram, Tamil Nadu. The specific geographic location was identified by coordinates 11°23′23.1″N/79°43′05.3″E. Healthy seeds of *Vigna radiata* L. (Co-6) variety were used for the research, were surface sterilized with





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0.2% mercuric chloride(HgCl₂) solution for 2 minutes and washed extensively with sterile double distilled water (ddH₂o). *Vigna radiata* L.Seeds were distributed and sown in 90 pots which were subsequently separated into six groups. Each pot was filled with a mixture of Red soil, sand, and farm yard manure in a ratio of (1:1:1). Plants were treated in Control, NaCl (80mM) Alfa-Toc (200mg/L) Brassinolide (4mg/L), NaCl (80mM) + Alfa-Toc(200mg/L), and NaCl (80mM) + BL (mg/L). To maintain particular salinity levels in pots, the soil samples from each pot were checked with Electrical Conductivity Meter at regular intervals. Plants were harvested for analyzing morphological and chlorophyll pigment analysis at the 25th, 35th, and45th, days after sowing [DAS].

Root length

Root length was measured from the point where the root and shoot meet to the tap root. The length of the lateral roots was taken together to get the total length of the root. The length of the roots was measured in cm plant -1.

Stem length

The length of the stem was measured from the tip to the point where the stem meets the root. And The length of the stems is given in cm plant ⁻¹.

Fresh weight

The roots and shoots of the plant were washed and cleaned with tap water, and gently dried with tissue paper. After drying automatic balance (Model: XK3190-A7M) was used to measure the fresh weights of the roots and shoots. The values were measured and expressed in gm plant ⁻¹.

Dry weight

After getting the fresh weight of plants, they were dried in a hot air oven at 60°C for 48 hours. After the plants were dry, their weights were measured, and they were kept in the same oven until their dry weights were the same. The values were expressed in gm plant ⁻¹.

Leaf Area

The total leaf area was measured using LICOR Photo Electric area meter (Model L1-3100 Lincoln, USA.) and expressed in cm² plant ⁻¹.

Chlorophyll pigment and carotenoid contents

Fresh leaf samples of 500 mg were taken from the plants and crushed with a pestle and mortar added 10 ml of 80% acetone. The mixture was then centrifuged at 800 rpm for 15 minutes. The extraction process was repeated, and the upper supernatant was collected, and mixed, with 20 ml of acetone to make a final volume of 20 ml. The spectrophotometer was used to measure absorption at 645, 663, and 480, nm against a blank of 80% acetone. The chlorophyll and carotenoids in the leaves were taken out and their amounts were measured using Arnon's method (1949). The findings are given in mg/gram of fresh weight.

RESULT AND DISCUSSION

Root length.

NaCl in *Vigna radiata* L. causes the stress declining the growth. In this experimental study, it was found that the plants treated with NaCl showed much decline in root length growth as compared to the control. However, plants treated with Alfa-Toc and Brassinolide show enhanced root growth than NaCl and control-treated plants. NaCl + BL and NaCl + Alfa-Toc treated plants also show increased root length than NaCl treated plants. (Fig-1)Plant development is negatively impacted by salinity due to low osmotic potential in soil solutions and nutritional imbalances. [26] Salinity causes regulated biophysical constraints on cell growth, which hinder root uptake and development. [27]. Other plants also show decreased root length with an increase in salinity like Tomato [28]*Brassica napus* [29]foliar application of Alfa-toc leads to notable increases in the growth of roots in sunflower *Helianthus*





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annuus L. [30].and *Glycine max* [31]Applying brassinolide via foliar spray notably countered the detrimental effects of salt stress, fostering increased root length in both wheat and *Vigna radiata* L.[32,33].

Stem length.

Salt stress hampers the growth and development of plants by restricting cell division and expansion, which hampers photosynthesis and results in shortened stem.[34] As per our results, the stem length of plants treated with NaCl declined much compared to the control, while the plants treated with Alfa-Toc and BL showed increased stem length compared to the control, the plants treated with Alfa-Toc+ NaCl and BL + NaCl showed enhanced growth than plants treated with NaCl alone. (Fig-2) High salinity levels have a substantial negative effect on plant growth by inducing osmotic stress and disrupting the balance of ions and nutrients. These disruptions have adverse consequences on the plant's essential physiological and biochemical processes necessary for its overall well-being and development.[35].Salt levels have been associated with a decrease in the growth of stem in sugar beet (*Beta vulgaris* L.) [36].and a similar outcome has been observed in sorghum[37]. Foliar applications of α -tocopherol increased growth traits such as stem length and ameliorated the salinity stress effects in faba beans [38].Alphatocopherol rejuvenates plant growth by neutralizing singlet oxygen species, acting as a shield for photosystems, particularly in saline environments. [39]. Brassinosteroids (BRs) are commonly occurring plant hormones crucial for both plant growth and managing stress [40]. Application of brassinolide (BL) treatments prompts elongation, cell division, and differentiation, ultimately supporting the growth of plants. [41]BL application has been associated with an increase in stem length in both sunflower [42] and cowpea.[43].

Fresh weight and dry weight.

The fresh weight and dry weight decrease under the saline treatments. In this study, the plants treated with NaCl show a drastic decline in weight compared to the control. In comparison, the plants treated with Alfa-Toc and BL show increased fresh and dry weights. However, plants treated with Alfa-Toc + NaCl and BL + NaCl show increased growth than NaCl-treated plants.(Fig-3,4). The application of NaCl treatment showed adverse effects on both the fresh weight and dry weight of *Brassica juncea*. [44]. *Lactuca sativa* [45] and Sorghum plants [46]. Our results agree with the results found in *Zea mays* [47], and wheat [48], and many others who observed the marked decreasein all the growth parameters. Application of Alfa-tocopherol at a concentration of 200mg/L-1 via foliar spray boosted both the fresh and dry weight of *Vigna radiata* L. [49] Alfa-toc also increased the fresh and dry weight of plants like carrot (*Daucus carota* L.). [50] *Glycine max* L. [51] and *Safflower* [52]The effects of salt are mitigated and the fresh and dry weight is increased by brassinolide foliar spray in *barley* [53]The treatment resulted in enhanced growth and higher dry and fresh weights in both *Medicagosativa* L. (alfalfa) and *Brassica juncea* plants. [54,55]and *Triticum aestivum* L. [56].

Leaf Area.

Leaf Area decreased inthe *Vigna radiata* L. in NaCl-treated plants when compared to the control, However, plants treated with Alfa-Toc and BL have found increased Leaf- Area compared to the control, plants treated with Alfa-Toc + NaCl and Bl + NaCl show increased leaf area than NaCl treated plants but lesser than control plants. (Fig-5)Many types of plants have shown that salt significantly affects leaf area. [57]. with increasing salt stress decreased leaf-Area has been found in many plants Like, sugar beet (*Beta vulgaris* L.) [58].Rice (*Oryza sativa*) [59]. Barley (*Hordeum vulgare* L.) [60]. The application of Alfa-Tocopherol through foliar has been observed to alleviate the effects caused by salt stress and enhance growth parameters, like leaf area, in *Helianthus annuus*L. [61]. Enhanced leaf area by Alfa-Toc application has been also found in *Vicia faba*L. [62]Application of Brassinolide externally has been demonstrated to boost both the height of the plant and the area of its leaves in maize, and same plant [63,64] and helps in mitigating the salt stress effects on plants.

Chlorophyll pigments.

Salinity causes a reduction in photosynthetic pigment concentrations due to a reduction in water potential. NaCl prevents the uptake of Nitrogen which is an important constituent of Chlorophyll. [65]. Declining the content of Chlorophyll, a and b, and total chlorophyll in *Vicia faba*, and *Vigna unguiculata* L. [66,67]. In our experimental study,





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the chlorophyll content of plants treated with NaCl showed a decline compared to the control, However, plants treated with Alfa-Toc and BL show enhanced chlorophyll content compared to control, and the plants treated with Alfa-toc+ NaCl and BL+ NaCl show a bit increased content than NaCl Treated plants. (Fig-6,7,8) our results are in accordance with the results found in *Hibiscus rosa sinensis* L[68]. Zea mays [69]. Oryza sativa. [70] Vigna radiata L. [71]. Alfa-Toc and Brassinolide assists in reducing the harmful impact of salt stress and improving the levels of photosynthetic pigments.

Carotenoid content

Carotenoids are supposed to be helpful in protection of the photosynthetic system of those plants that generally grown under extreme environmental conditions by enhancing the process of ROS scavenging.[72]. As per our results, the plants treated with Alfa-Toc and BL have high carotenoid content compared to the control, however, plants treated with NaCl have less carotenoid content compared to control, and the plants treated with Alfa-Toc + NaCl and BL + NaCl have high carotenoid content than NaCl treated plants. (Fig-9) Brassinolide application enhances the content of carotenoids in plant leaves [73]Carotenoids can deactivate singlet oxygen by absorbing it, causing them to enter a triplet state that dissipates through thermal decay, effectively neutralizing its effects. [74]Alfa-Toc plays a significant role in plants by increasing the carotenoid content, as observed in *Triticum aestivum*. [75].

CONCLUSION

High salt levels pose a substantial environmental challenge, particularly constraining plant productivity, especially in dry and semi-dry climates. In the present study, NaCl causes negative impacts on physiological, biochemical, and pigment composition in *Vigna radiata* L. (Co-6) variety. Application of brassinolide and Alfa-tocopherol helps in mitigating the negative impacts of salinity stress, and plays an eminent role in enhancing growth, development, and pigment composition in green gram(*Vigna radiata* L. Co-6 variety)Therefore, spraying these two Plant growth regulators might be a suitable approach in Ameliorating the salt stress in *Vigna radiata* L and in enhancing biomass and pigment composition.

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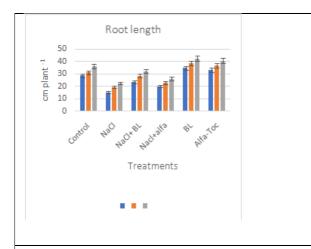


Fig. 1. Effect of exogenous application of Alfa-Toc and BL on Root length of Vigna radiata L. [CO-6 variety] under 80mM NaCl stress. Values represented in Bars are the mean of three replicates [n=3] and [±] standard error.

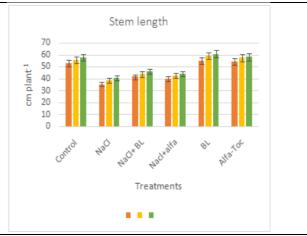
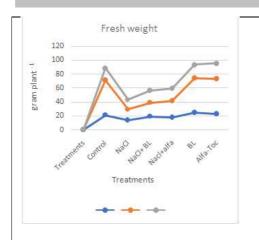


Fig. 2. Effect of exogenous application of Alfa-Toc and BL on stem length of *Vigna radiata* L. [CO-6 variety] under 80mM NaCl stress. Values represented in Bars are the mean of three replicates [n=3] and [±] standard error.







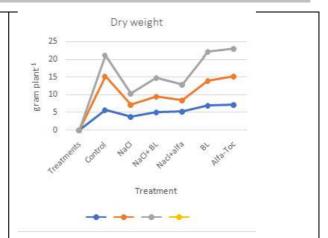
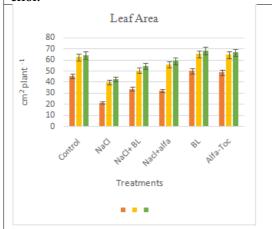


Fig. 3. Effect of exogenous application of Alfa-Toc and BL on Fresh weight of Vigna radiata L. [CO-6 variety] under 80mM NaCl stress. Values represented in Bars are mean of three replicates [n=3] and [±] standard error.

Fig. 4. Effect of exogenous application of Alfa-Toc and BL on Dry weight of Vigna radiata L. [CO-6 variety] under 80mM NaCl stress. Values represented in Bars are mean of three replicates [n=3] and [±] standard error.



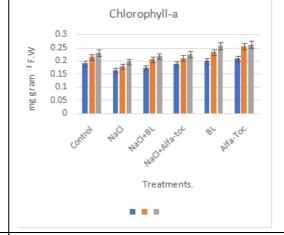
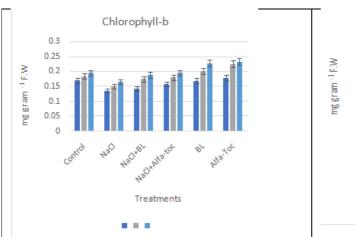


Fig. 5. Effect of exogenous application of Alfa-Toc and BL on leaf area of *Vigna radiata* L. [CO-6 variety] under 80mM NaCl stress. Values represented in Bars are the mean of three replicates [n=3] and [±] standard error.

Fig. 6. Effect of exogenous application of Alfa-Toc and BL on Chlorophyll-a of *Vigna radiata* L. [CO-6 variety] under 80mM NaCl stress. Values represented in Bars are the mean of three replicates [n=3] and [±] standard error.







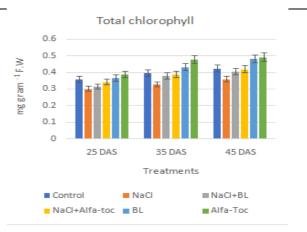


Fig. 7. Effect of exogenous application of Alfa-Toc and BL on Chlorophyll-b of *Vigna radiata* L. [CO-6 variety] under 80mM NaCl stress. Values represented in Bars are the mean of three replicates [n=3] and [±] standard error.

Fig. 8. Effect of exogenous application of Alfa-Toc and BL on Total Chlorophyllof *Vigna radiata* L. [CO-6 variety] under 80mM NaCl stress. Values represented in Bars are the mean of three replicates [n=3] and [±] standard error.

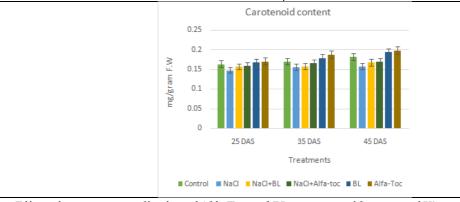


Fig. 9. Effect of exogenous application of Alfa-Toc and BL on carotenoid content of *Vigna radiata* L. [CO-6 variety] under 80mM NaCl stress. Values represented in Bars are the mean of three replicates [n=3] and [±] standard error.

